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Epilepsy-associated genes: an update

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ABSTRACT

Purpose: To provide an updated list of epilepsy-associated genes based on clinical-genetic evidence. Methods: Epilepsy-associated genes were systematically searched and cross-checked from the OMIM, HGMD, and PubMed databases up to July 2023. To facilitate the reference for the epilepsy-associated genes that are potentially common in clinical practice, the epilepsy-associated genes were ranked by the mutation number in the HGMD database and by case number in the China Epilepsy Gene 1.0 project, which targeted common onlines.

Results: Based on the OMIM database, 1506 genes were identified to be associated with epilepsy and were classified into three categories according to their potential association with epilepsy or other abnormal phenotypes, including 168 epilepsy genes that were associated with epilepsies as pure or core symptoms, 364 genes that were associated with neurodevelopmental disorders as the main symptom and epilepsy, and 974 epilepsy-related genes that were associated with gross physical/systemic abnormalities accompanied by epilepsy/seizures. Among the epilepsy genes, 115 genes (68.5%) were associated with epileptic encephalopathy. After cross-checking with the HGMD and PubMed databases, an additional 1440 genes were listed as potential epilepsy-associated genes, of which 278 genes have been repeatedly identified variants in patients with epilepsy. The top 100 frequently reported/identified epilepsy-associated genes from the HGMD database and the China Epilepsy Gene 1.0 project were listed, among which 40 genes were identical in both sources.

Significance: Recognition of epilepsy-associated genes will facilitate genetic screening strategies and be helpful for precise molecular diagnosis and treatment of epilepsy in clinical practice.

1. Introduction

The widespread applications of next-generation sequencing technologies have accelerated the discovery of genes associated with epilepsy. Genetic factors are believed to be the major contributors to the cause of epilepsy in up to 80% of people with epilepsy [1–3]. In 2017, we summarized 977 genes that are associated with epilepsy based on clinical-genetic evidence [4]. Over the past six years, a great number of novel epilepsy-associated genes have been reported, which has led to encouraging progress in the field of epilepsy genetics. Recently, two genetic databases for epilepsy have been established: the Seizure-Associated Genes Across Species (SAGAS) database (www.

sagas.ac), which included 2876 epilepsy-associated genes by gathering the genes potentially associated with seizures or epilepsies in multiple species [5], and the Gene4epilepsy database (https://github.com/bahlol ab/genes4epilepsy), which contained 926 genes by comparing genes included on the epilepsy panels of four clinical diagnostic providers and two research resources, in which genes were mostly associated with developmental and epileptic encephalopathies [6]. From the perspective of clinical practice, it is important to know which genes are clinically associated with epilepsy and the relationships between epilepsy and the other abnormal phenotypes. As the pace of epilepsy-associated gene discovery accelerates, we now update the list of epilepsy-associated genes based on the genes that are currently reported to be related to

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epilepsy clinically. We searched for epilepsy-associated genes by reviewing the literature in the Online Mendelian Inheritance in Man (OMIM) database, Human Gene Mutation Database (HGMD), and PubMed. The purpose of the present study is to provide a list of epilepsy-associated genes that is potentially useful for facilitating the application of genetic tests in the clinical genetic diagnosis and precise treatment of epilepsy.

2. Methods

To update the list of genes associated with epilepsy, a systematic review was conducted based on data from online databases until July 2023. Following the methods described in the first version [4], genes were first searched in the OMIM database (https://www.omim.org/) using the following terms: "epilepsy/epilepsies/epileptic", "seizure/seizures", and "convulsion/convulsions". The OMIM database is the primary repository of comprehensive and curated information on genes and genetic phenotypes and the relationships between them [7]. Over 4000 items were retrieved from the OMIM database, among which those related to epilepsy with well-characterized genetic mutations in humans were included. The excluded items were (1) that in clinical genetic reports (studies) without definite molecular genetic confirmation; (2) that from linkage studies without identification of specific genes; (3) that were associated genomic rearrangement without definite causative genes; (4) the studies in which the patients with epilepsy had not been subjected to genetic tests; and (5) genes related to epilepsy in animal models but not in humans yet. The 977 genes in the first version were rereviewed and reclassified. Some genes have been upgraded with novel epilepsy-related evidence, whereas some were removed from the list of epilepsy-associated genes. Finally, 1506 genes were identified to be associated with epilepsy, including 680 previously reported in the first version and 826 newly obtained genes. According to the phenotype described currently in the OMIM database, these genes were classified into three categories: (1) epilepsy genes, i.e., genes that were potentially associated with epilepsies as pure or core symptoms, including genes that may be associated with multiple phenotypes other than epilepsy or seizures but have also been reported in cases presented pure epilepsy or seizures; (2) neurodevelopment-associated epilepsy genes, i.e., genes that were associated with gross brain developmental malformations or neurodevelopmental disorders that were accompanied by epilepsy or seizures; and (3) epilepsy-related genes, i.e., genes associated with gross physical or other systemic abnormalities and accompanied by epilepsy or seizures.

Then, genes with mutations in the epilepsy-related phenotypes were comprehensively searched in the HGMD (http://www.hgmd.cf.ac.uk/ ac/index.php, version: HGMD Professional 2022.4). The HGMD constitutes a comprehensive collection of published germline mutations in nuclear genes that are thought to underlie or are closely associated with human inherited disease [8]. Furthermore, epilepsy-related publications from January 2017 to July 2023 were searched in the PubMed database. The terms "epilepsy/epilepsies/epileptic", "seizure/seizures", and "convulsion/convulsions" were used for the search. The results from these databases were cross-checked. The genes that were reported to be associated with epileptic phenotypes in the HGMD and/or PubMed databases but were not yet included in the OMIM database were listed as potential epilepsy-associated genes, which included 1367 genes from HGMD and an additional 73 genes that appeared only in PubMed. Compared with the previous version from 2017, we excluded 46 potential epilepsy-associated genes from the EpilepsyGene databases (http ://61.152.91.49/EpilepsyGene/) due to insufficient evidence.

Genes from the two online databases for epilepsy, the SAGAS database and the Genes4Epilepsy database, were reviewed and checked by gene ID. There was duplicate recruitment of synonymous genes in the SAGAS database, including ADGRV1/ADRV1, AKAP5/AKAP150, CDKL5/CDLK5, COX2/MTCO2, DNAJC21/DNAJA5, DNM1/DMN1, IFIH1/IFIHI1, MIR128/MIR128-1, MTTK/MT-TK, MTTW/MT-TW,

MTOR/MTORC, MECP/MECP2, OCRL/OCRL1, PIK3CA/PI3KCA, SRP54/SCN8, SLC35A2/SCL35A2, SIK1/SIK1B, and SAMD12/FCMTE1. The duplicate recruiting of synonymous genes in the Genes4Epilepsy database were ADPRHL2/ADPRS, H3-3A/H3F3A, H3-3B/H3F3B, and PLPBP/PROSC. After excluding duplicates, the epilepsy-associated genes were confirmed to be 2858 in the SAGAS database (2876 genes in the previous report) and 922 in the Genes4Epilepsy database (926 genes in the previous report), which were compared with the updated epilepsy-associated genes in the present study (n = 2946).

To facilitate the reference for the epilepsy-associated genes that are potentially common in clinical practice, we ranked the epilepsyassociated genes by the number of mutations associated with epileptic phenotypes in the HGMD database. Similarly, we ranked the causative genes by case number using data from the China Epilepsy Gene 1.0 project, which were obtained from trio-based whole-exome sequencing (WES) analysis of 2757 unrelated cases with common epilepsies, reflecting the scenery of the real world. Trio-based WES was performed in all cases. The included variants met the criteria in two aspects. First, the variants were *de novo*, recessive from the two asymptomatic parents each, hemizygous, or cosegregated variants, which highlights the genetic difference between the affected child and the parents and thus explains the occurrence of phenotype in a given family (trio). Second, the variants met the following criteria on the minor allele frequency (MAF): (1) the MAF for de novo variants is < 0.0001, according to gnomAD in the general population or East Asian population; (2) homozygotes/hemizygotes are not present in the controls, according to gnomAD; (3) the MAF of biallelic variants of compound heterozygotes is < 0.005 for each heterozygous variant; and (4) the MAF of cosegregated variants is < 0.005 and at least two affected individuals in a family. The top 100 epilepsy-associated genes from the HGMD database and the China Epilepsy Gene 1.0 project were listed.

3. Results

The present study established a list of 2946 epilepsy-associated genes (Fig. 1A) and grouped the genes according to their potential association with epilepsy and the relationship between epilepsy and other clinical abnormalities. These genes included 168 epilepsy genes, 364 neurodevelopment-associated epilepsy genes, and 974 epilepsy-related genes, which were recruited in OMIM (Fig. 1B, 1506 in total). An additional 1440 genes were retrieved from HGMD and PubMed and were listed as potential epilepsy-associated genes, among which variants in 278 genes have been repetitively identified in patients with epilepsy. Therefore, based on clinical-genetic evidence currently available, a total of 1784 (1506 \pm 278) genes are potentially significant in practice.

Compared with the SAGAS database and the Genes4Epilepsy database, 807 genes were common among the three databases. There were 62 genes from the Genes4Epilepsy database but were not enrolled in the present epilepsy-associated gene list, which were mostly genes involved in development but without clinical evidence on association with epilepsy, e.g., AKT1, which was associated with schizophrenia and development disorders [9,10], and ZMYND8, which was associated with autism spectrum disorder [11]. Almost 1000 genes from the SAGAS database were not included in the present list due to insufficient clinical evidence, such as ADAM10 [12] and HAS3 [13], which had only epilepsy-related evidence from animal studies. A total of 1025 genes uniquely appeared in the present list of epilepsy-associated genes (Fig. 1C).

3.1. Epilepsy genes

Overall, 168 genes were classified as epilepsy genes, including 82 genes from the previous version and 86 newly reported genes. Two epilepsy genes in the previous list, *ADRA2B* and *SCN9A*, were downgraded to potential epilepsy-associated genes due to the lack of definite evidence, such as a lack of cosegregation studies, high frequencies of

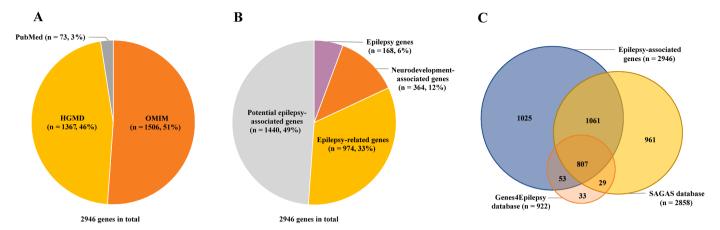


Fig. 1. The overall illustration of epilepsy-associated genes.

- (A) Epilepsy-associated genes identified in the OMIM, HGMD, and PubMed databases.
- (B) The epilepsy-associated genes were classified into four categories according to their potential association with epilepsy.
- (C) Comparison of genes in the SAGAS database, the Genes4Epilepsy database, and the updated epilepsy-associated genes of the present study.

variants in the general population, and additional causative factors in the cases [14,15].

These 168 genes were listed according to the clinical features of epileptic phenotypes (Table 1). The majority of epilepsy genes (90, 53.6%) were of autosomal dominant inheritance; 62 genes were of autosomal recessive inheritance; 4 genes were of both autosomal dominant and recessive inheritance; 9 genes were of X-linked inheritance; and 3 genes had undefined inheritance pattern. Among these genes, 68.5% (115/168) were causative genes of epileptic encephalopathy, and 13 genes (7.7%) were causative genes of progressive myoclonic epilepsy. Heterogeneity in phenotypes and inheritance patterns has been observed in several genes, such as the *SLC12A5* variants that were associated with epileptic encephalopathy in autosomal recessive inheritance but with idiopathic generalized epilepsy in autosomal dominant inheritance [16–18].

3.2. Neurodevelopment-associated epilepsy genes

A total of 364 genes were categorized as neurodevelopment-associated epilepsy genes, which was 291 more than that in the first version. These genes were listed according to phenotype features (Table 2), which cover a wide range of brain abnormalities from focal brain malformations (e.g., cortical dysplasia) to general brain malformations (e.g., microcephaly) and others.

The neurodevelopment-associated genes presented recessive inheritance in the majority (214/364, 58.8%), including 206 genes with autosomal recessive inheritance and 8 genes with X-linked recessive inheritance. Four genes were associated with the inheritance of both autosomal dominant and recessive features, including *AP1G1*, *ATAD3A*, *CLCN3*, and *NARS1*.

An increasing number (211) of genes associated with general neurodevelopmental disorders have been identified, including 49 genes with multisystemic abnormalities (e.g., cardiac defects, renal anomalies, and gastrointestinal issues). For instance, variants in *ADNP* cause Helsmoortel-Van der Aa syndrome (HVDAS), which is a neurodevelopmental disorder characterized by impaired intellectual development/motor delay, autism spectrum disorder, facial dysmorphisms, hypotonia, congenital heart disease, visual difficulties, and gastrointestinal issues [19]. In contrast, 162 genes have not been associated with multisystemic abnormalities reported thus far.

3.3. Epilepsy-related genes

Up to 974 genes were categorized as epilepsy-related genes, which was 438 more than that in the first version. These genes are listed in

Table 3, with the detailed epilepsy-related phenotypes in Table S1.

Genes with recessive inheritance accounted for the majority (665/974, 68.3%), which included 605 genes with autosomal recessive inheritance and 60 genes with X-linked recessive inheritance. Both autosomal dominant and recessive inheritance features were observed in 26 genes.

Variants in these genes result in diseases presenting gross physical or other systemic abnormalities that are accompanied by epilepsy or seizures, e.g., mitochondrial complex deficiency, cardiac disease, and metabolic disorders. Cardiac diseases, such as long QT syndrome, congenital heart defects, cardiomyopathy, and heart block, potentially present cardiac arrhythmias and subsequently recurrent syncope, seizures, and even sudden death [20]. Metabolic disorders, typically as combined oxidative phosphorylation deficiency and glycogen storage disease, are potentially associated with seizures due to the involvement of the central nervous system.

3.4. Potential epilepsy-associated genes

In total, 1440 genes were classified as potential epilepsy-associated genes, including 1367 genes from the HGMD database and an additional 73 genes from PubMed (Table 4). The reported epileptic phenotypes with the corresponding number of mutations in each gene are listed in Table S2. The majority of the genes (1162) had only a single variant identified in epileptic phenotypes and thus warrant further studies to validate their association with epilepsy. In contrast, variants of 278 genes have been repeatedly identified in patients with epilepsy, including 34 genes with 5 or more variants identified in epileptic phenotypes. Validating experimental evidence supporting the gene-disease association was also obtained for some genes, such as *PRICKLE2* [21], *KCNH5* [22], *GABRA6* [23], *KCND2* [24,25], and *UNC13B* [26], which are potential epilepsy genes but have not yet been included in OMIM.

3.5. Frequently reported/identified epilepsy-associated genes

The top 100 epilepsy-associated genes obtained from the HGMD and the China Epilepsy Gene 1.0 project are summarized in Table 5. SCN1A is the most frequent epilepsy gene in both databases. While 40 genes appeared identically, the epilepsy-associated genes differed in 60 genes in the two databases, potentially due to the study subjects and publication bias. For example, STXBP1 has been associated with rare and severe epileptic encephalopathy and is ranked seventh among the top 100 epilepsy-associated genes from the HGMD database, potentially due to the severe phenotype that is commonly the subject of studies; however, STXBP1 did not appear in the top 100 epilepsy-associated genes

Table 1 Epilepsy genes (n = 168).

Epilepsy genes ($n = 16$	8).		
Phenotype (in order of the onset age)	Inheritance	Genes by 2017	Genes updated
Neonatal			
Pyridoxine-	AR	ALDH7A1, PNPO	PLPBP
dependent epilepsy		. , .	
Benign familial	AD	KCNQ2, KCNQ3	
neonatal seizures			
(BFNS)			
Myoclonus,	AD		KIF5A
intractable,			
neonatal			
Rigidity and	AR		BRAT1
multifocal seizure			
syndrome, lethal			
neonatal			
Infantile and			
childhood			
Familial infantile	AR	TBC1D24	
myoclonic epilepsy			
(FTLE)			
Benign familial	AD	PRRT2, SCN2A,	
infantile seizures		SCN8A	
(BFIS)			
Epileptic	AD	CACNA1A, CHD2,	AP2M1, ATP1A2,
encephalopathy	112	DNM1, EEF1A2,	ATP1A3,
		FGF12, GABRA1,	ATP6V0A1,
		GABRB1,	ATP6V1A,
		GABRB3,	ATP6V0C,
		GABRG2, GNAO1,	CACNA1E, CDK19,
		GRIN2B, GRIN2D,	CELF2, CUX2,
		HCN1, KCNA2,	CYFIP2, FBXO28,
		KCNB1, KCNQ2,	FZR1, GABBR2,
		KCNT1, SCN1A,	GABRA2, GABRA5,
		SCN2A, SCN8A,	GABRB2, HNRNPU,
		SIK1, SLC1A2,	KCNC2, KCNT2,
		SPTAN1	MAST3, NEUROD2,
		SF IAIVI	
			NSF, NTRK2, NUS1,
			PACS2, PHACTR1,
			PPP3CA, RHOBTB2,
			RNF13, SCN3A,
			SETD1A, SETD1B,
			YWHAG
	AR	AARS1, ARV1,	ACTL6B, ADAM22,
		DOCK7, FRRS1L,	AP3B2,
		GUF1, ITPA,	CACNA2D1, CAD,
		NECAP1, PLCB1,	CNPY3, CPLX1,
		SCN1B, SLC12A5,	DALRD3,
		SLC13A5,	DENND5A, DMXL2,
		SLC25A12,	GAD1, GLS, GOT2,
		SLC25A22,	GRIN1, HID1,
		ST3GAL3,	MDH1, MDH2,
		ST3GAL5, SZT2,	NAPB, NRROS,
		TBC1D24, UBA5,	PARS2, PIGB, PIGP,
		WWOX	PIGS, SLC38A3,
			SYNJ1, TRAK1,
			UFSP2, UGDH,
			UGP2
	AD, AR	STXBP1	
	XLD	CDKL5	SLC35A2, SMC1A
	XLR		ARX
	XL	ALG13, ARHGEF9,	FGF13, GABRA3
		PCDH19	
Dravet syndrome	AD	SCN1A	
(DS)			
Familial febrile	AD	ADGRV1,	HCN2
seizures (FFS)		GABRG2, SCN1A	
·/	AR	CPA6	
Generalized	AD	GABRD, GABRG2,	HCN2
epilepsy with	_	SCN1A, HCN1,	
febrile seizures plus		SCN1B, STX1B	
(GEFS+)		JULIE, JIAID	
Myoclonic-atonic	AD	SLC6A1	
epilepsy (MAE)	AD	OLCONI	
Focal epilepsy and	AD	GRIN2A	
speech disorder	AD	GIUIVZA	
speccii disorder			

Table 1 (continued)

Phenotype (in order of the onset age)	Inheritance	Genes by 2017	Genes updated
(FESD) with or without impaired intellectual development			
Childhood absence	AD	GABRG2	
epilepsy (CAE)	UN	CACNA1H, GABRA1, GABRB3	
Juvenile and later			
Juvenile absence epilepsy (JAE)	AD	CLCN2, EFHC1	
Juvenile myoclonic epilepsy (JME)	AD	CACNB4, CLCN2, EFHC1, GABRD	ICK
	UN	GABRA1	
Idiopathic	AD	CACNB4, CLCN2,	RORB, HCN2 ,
generalized		GABRD, KCNMA1,	HCN4
epilepsy (IGE)	****	SLC12A5, SLC2A1	
m 11 1 1 1	UN	CACNA1H, CASR	MIDOCA DADGEE
Familial adult myoclonic epilepsy (FAME)	AD		TNRC6A, RAPGEF2 YEATS2, SAMD12, STARD7, MARCH6
(ITHILL)	AR	CNTN2	biriid, minterio
Familial temporal lobe epilepsy (FTLE)	AD	CPA6, GAL, LGI1	RELN
Not specific	AD	WONG1	CEMA CD
Progressive myoclonic epilepsy (PME)	AD AR	KCNC1 CERS1, CSTB, EPM2A, GOSR2, KCTD7, LMNB2, NHLRC1, PRDM8, PRICKLE1, SCARB2	SEMA6B SLC7A6OS
Nocturnal frontal lobe epilepsy	AD	CHRNA2, CHRNA4, KCNT1	
(NFLE)	UN	CHRNB2	
Familial focal epilepsy with variable foci (FFEVF)	AD	DEPDC5	NPRL2, NPRL3, SCN3A

Bold italics, with multiple epilepsy phenotypes.

AD, autosomal dominant; AR, autosomal recessive; UN, unknown; XL, X-linked; XLD, X-linked dominant; XLR, X-linked recessive.

from the China Epilepsy Gene 1.0 project, which targeted common epilepsies, including common epilepsy with favorable outcomes (e.g., *UNC13B*). This explains the appearance of several newly identified epilepsy genes, such as *HCFC1* [27], *BCOR* [28], *PKD1* [29], *AFF2* [30], *MED12* [31], *NEXMIF* [32], *BSN* [33], *CELSR1* [34], *FAT1* [35], *LAMA5* [36], *BRWD3* [37], *CELSR3* [38], *FRMPD4* [39], *RYR2* [40], *SHROOM4* [41], *UNC13B* [26], *APC2* [42], *CHD4* [43], *PGM3* [44], *MPDZ* [45], *ATP6VOC* [46] and *UNC79* [47], in the epilepsy-associated genes from the China Epilepsy Gene 1.0 project.

4. Discussion

The gene-disease association refers to three levels in general: (1) what phenotype a gene is associated with; (2) whether a gene is associated with the phenotype (pathogenic potential of a gene); and (3) how a gene is associated with the phenotype (pathogenic feature of a gene). Evaluating gene-disease associations is a complex task that involves multiple aspects of evidence. The present study focused on the first level of gene-disease association, i.e., which gene is potentially associated with epilepsy and the relationship between epilepsy and other clinical abnormalities. A list of epilepsy-associated genes was established based on clinical-genetic evidence, which provides an overview of the epilepsy-associated genes and is potentially helpful for gene screening in clinical practice. The epilepsy-associated genes were further classified into four categories, which are potentially useful for clinical genetic diagnosis and precise treatment.

Table 2 Neurodevelopment-associated epilepsy genes (n = 364).

Table 2 (continued)

Phenotype	Inheritance	Genec by 2017	Genes undated	Phenotype	Inheritance	Genes by 2017	Genes updated
Phenotype Focal or multifocal brain	Inheritance	Genes by 2017	Genes updated				HEPACAM, HSPD1, HYCC1,
malformation grey matter							MLC1, KARS1, POLR3A,
Holoprosencephaly	AD	PTCH1	CNOT1				POLR3B,
Trotoprosencephary	AR	110111	PLCH1				POLR3K, PYCR2
Pseudo-TORCH	AR	OCLN					RARS1,
syndrome							RNASET2,
Polymicrogyria	AR	ADGRG1, FIG4	COL3A1, PI4KA				SNORD118,
	UN	ADGRG1					UFM1, VPS11
CK syndrome	XLR	NSDHL		General brain	XLR		PLP1
Megalencephaly-	AD	PIK3R2	AKT3, CCND2	malformation & others			
polymicrogyria-				Lissencephaly	AD	TUBA1A,	CEP85L, MACF1
polydactyly- hydrocephalus syndrome				Lisscheephary	71D	PAFAH1B1	GLI OSL, WITGIT
Rolandic epilepsy,	UN	SRPX2			AR	CDK5, KATNB1,	TMTC3
impaired intellectual	011	514112				LAMB1, NDE1	
development, and speech					XL	DCX	
dyspraxia				Mega-corpus-callosum	AD		MAST1
Periventricular	AD	ERMARD	ARF1, MAP1B,	syndrome with cerebellar			
heterotopia			NEDD4L	hypoplasia and cortical			
	AR	ARFGEF2		malformations			
	XLD	FLNA		Microcephaly with	AD	KIF11	D
Band heterotopia	AR	DADATES.	EML1	variable abnormalities	AR	DIAPH1,	DTYMK, SLC1A4
Subcortical laminar	AD	PAFAH1B1				IER3IP1, MED17, PNKP,	TUBGCP6, TUBGCP2,
heterotopia	XL UN	DCX	MTOR			PPP1R15B,	WDR4, YIPF5
Focal cortical dysplasia, type II, somatic	UN		MIOR			QARS1, RTTN,	WDIG, 11113
Complex cortical	AD	KIF2A, KIF5C,				STAMBP,	
dysplasia with other brain	1110	TUBB2A,				TRMT10A	
malformations		TUBB2B,		Primary microcephaly	AD		LMNB1
		TUBB3, TUBG1		1 7	AR	ANKLE2, ASPM,	CIT, NCAPD3,
	AR	•	APC2, CAMSAP1,			CENPE, CENPJ,	PDCD6IP, STIL,
			CTNNA2			WDR62, SASS6	ZNF335
Pitt-Hopkins like	AR	CNTNAP2		Macrocephaly,	AR	HERC1	
syndrome				dysmorphic facies, and			
Occipital cortical	AR	LAMC3		psychomotor retardation			
malformations				Polyhydramnios,	AR	STRADA	
Pontocerebellar	AR	AMPD2, CLP1,	ATAD3A,	megalencephaly, and			
hypoplasia		EXOSC3, PCLO,	CDC40, EXOSC9,	symptomatic epilepsy	AD		NALON TROK
		SEPSECS,	MINPP1, PPIL1,	Hypotonia with psychomotor retardation	AR		NALCN, TBCK, UNC80
		TSEN2, TSEN15,	PRDM13, RARS2, SLC25A46,	and characteristic facies			014000
		TSEN54, VPS53	TBC1D23, TOE1,	Hydrocephalus	AD		SMARCC1,
		1011101, 11000	VPS51	,			TRIM71
Cerebellar hypoplasia/	AR		CACNA2D2,		AR	CCDC88C,	
atrophy			EXOSC5, OXR1,			MPDZ	
1 3			VLDLR	Brain small vessel disease	AD	COL4A2	COL4A1
Diencephalic-	AR		PCDH12	Schizencephaly	UN	EMX2	SIX3
mesencephalic junction				Neurodevelopmental	AD		ADGRL1, AGO1,
mescricephane junetion							
dysplasia syndrome				disorder			ARFGEF1, BAP1,
dysplasia syndrome Dentatorubro-	AD	ATN1		disorder			BPTF, BRPF1,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy				disorder			BPTF, BRPF1, CACNA1C,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia	AD	ATN1 SLC20A2, XPR1		disorder			BPTF, BRPF1, CACNA1C, CACNA1I,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification			JAM2, MYORG	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter	AD AR		•	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus	AD AR AD	SLC20A2, XPR1	JAM2, MYORG CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter	AD AR AD AR		CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum	AD AR AD AR XLR	SLC20A2, XPR1	CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy,	AD AR AD AR	SLC20A2, XPR1	CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset,	AD AR AD AR XLR	SLC20A2, XPR1	CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and	AD AR AD AR XLR	SLC20A2, XPR1	CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum	AD AR AD AR XLR AR	SLC20A2, XPR1	CDH2	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and	AD AR AD AR XLR	SLC20A2, XPR1	CDH2 L1CAM TBCD	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1	CDH2 L1CAM TBCD ATP11A,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, H3-3B, H4C3, H4C5, HDAC4,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNAI1, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNAI1, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1 SLC12A6	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A AIMP1, AIMP2,	disorder			BPTF, BRPF1, CACNA1L, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL KAT8, KCNN2,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1 SLC12A6	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A AIMP1, AIMP2, ARSA, CNP,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL KAT8, KCNN2, KDM6B, KMT2C,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1 SLC12A6	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A AIMP1, AIMP2, ARSA, CNP, DARS2, DEGS1,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL KAT8, KCNN2, KDM6B, KMT2C, KMT2E,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1 SLC12A6	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A AIMP1, AIMP2, ARSA, CNP, DARS2, DEGS1, EIF2B1, EIF2B2,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL KAT8, KCNN2, KDM6B, KMT2C, KMT2E, LMBRD2,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1 SLC12A6	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A AIMP1, AIMP2, ARSA, CNP, DARS2, DEGS1, EIF2B1, EIF2B2, EIF2B3, EIF2B4,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL KAT8, KCNN2, KDM6B, KMT2C, KMT2E, LMBRD2, MAPK8IP3,
dysplasia syndrome Dentatorubro- pallidoluysian atrophy Idiopathic basal ganglia calcification White matter Agenesis of the corpus callosum Encephalopathy, progressive, early-onset, with brain atrophy and thin corpus callosum Leukodystrophy/	AD AR AD AR XLR AR	SLC20A2, XPR1 SLC12A6	CDH2 L1CAM TBCD ATP11A, CLDN11, CSF1R, EIF2AK2, FBP2, TMEM106B, TMEM163, TUBB4A AIMP1, AIMP2, ARSA, CNP, DARS2, DEGS1, EIF2B1, EIF2B2,	disorder			BPTF, BRPF1, CACNA1C, CACNA1I, CDC42BPB, CHAMP1, CHD1, CHD3, CHD5, CSNK2B, DHDDS DHX30, DLL1, DPYSL5, FBXW7, FOXG1, FOXP1, FRMD5, GNA11, GRIA2, GRIA4, GRIK2, H3-3A, H3-3B, H4C3, H4C5, HDAC4, HECW2, HNRPH1, HNRPR, IRF2BPL KAT8, KCNN2, KDM6B, KMT2C, KMT2E, LMBRD2,

Table 2 (continued)

Tubic 2 (continued)			
Phenotype	Inheritance	Genes by 2017	Genes updated
			NOVA2, NR4A2, PAK1, PHF21A, POGZ, POLR2A, POU3F3, PPP2CA, PRKAR1B, RAB11B, RALA, RFX7, RNF2, SOX5, TANC2, TCF20, TRIO, TRPM3, UBE3A, VAMP2, WASF1, WDR26, ZIC1, ZMIZ1, ZSWIM6
	AR	MFSD2A, SNIP1	ZMIZ1, ZSWIM6 ADARB1, ATP9A, BCAS3, CACNA1B, CHKA, CPSF3, CRADD, DEAF1, DHPS, DOHH, DYNC112, EMC10, ESAM, EXOC2, EXOC7, EXOC8, EXT2, GPT2, GRM7, GTPBP2, HECTD4, HPDL, INTS8, INTS11, LNPK, MADD, MED11, MED27, MTHFS, NAE1, NFASC, NRCAM, NSRP1, NTNG2, OGDHL, P4HTM, PCDHGC4, PDZD8, PGAP1, PIDD1, PIGG, PIGK, PIGU, PLAA, PLXNA1, PPFIBP1, PRUNE1, PTPN23, PUS3, RALGAPA1, RBL2, SHQ1, SMPD4, SPATA5L1, SPTBN4, SVBP, TAF8, TBC1D2B, THUMPD1, TIAM1, TMEM222, TMX2, TRAPPC10, TRAPPC6B, TTC5, UBE3C, UBE4A, UFC1, VARS1, WARS1, WDR45B, YIF1B, ZNF142, ZNF526,
	AD, AR		ZNF668, ZNHIT3 AP1G1, ATAD3A, CLCN3, NARS1
	XLR	OPHN1	HS6ST2, BCORL1, ZC4H2
	XLD XL	CASK SYN1	HNRNPH2 SLITRK2
Neurodevelopmental disorder with multisystemic abnormalities	AD	ATN1, TSC1, TSC2	ASXL2, ADNP, AGO2, CDK8, CREBBP, CSNK2A1, CUL3, DPH5, GATAD2B, GNB2, KAT5, MED12L, PSMD12, PURA,

Table 2 (continued)

Phenotype	Inheritance	Genes by 2017	Genes updated
	AR	SPATA5, WDR73	RAC3, SIAH1, SPEN, SUPT16H, TRRAP, USP7, WAC, ZMYM2, CAPN15, DHX37, FRA10AC1, GEMIN4, INTS1.
			KIAA1109, PGM2L1, PIGN, PIGQ, PIGT, SARS1, SEC31A, THOC6, TMEM147.
			TMEM94, UBR7, VPS50, WARS2
	XLD		TCEAL1
	XLR		PIGA
	XL		NAA10, HUWE1

Bold italics, with multiple epilepsy phenotypes. AD, autosomal dominant; AR, autosomal recessive; UN, unknown; XL, X-linked; XLD, X-linked dominant; XLR, X-linked recessive.

In this study, 168 genes were classified as epilepsy genes, which was twice as many as in the first version in 2017 [4]. Variants in these genes cause pure or relatively pure epilepsies or syndromes with epilepsy as the core syndrome. These genes probably play a major role in patients with epilepsy as the primary clinical manifestation. It is noteworthy that genes associated with neurodevelopmental abnormalities or other systemic abnormalities are also potentially associated with pure epilepsies, such as *MED12* [31], *RYR2* [40], and *PKD1* [29], as shown in recent studies.

The present list included 364 neurodevelopment-associated epilepsy genes, which were associated with general or focal brain structural abnormalities. The etiology of epilepsy is currently classified into structural, genetic, infectious, metabolic, immune, and unknown [48]. However, the growing number of genes potentially associated with brain structure abnormalities suggests attention to the underlying genetic etiology of brain structure disorders, which implies significance in clinical practice, such as presurgical evaluation.

Up to 974 genes were considered as epilepsy-related genes. Variants in these genes result in diseases presenting gross physical or other systemic abnormalities that are accompanied by epilepsy or seizures. The typical clinical features of gross physical or systemic abnormalities provide useful evidence for clinical and genetic diagnosis. A comprehensive systemic examination is required to detect specific systemic involvement, such as cardiac examination, metabolic screening, and hormone blood tests, depending on the specific genetic etiology. Accurate and early genetic diagnosis is potentially helpful in the precise management of patients, such as riboflavin in patients with glutaric acidemia caused by *ETFDH* mutations [49].

In the present study, 1440 genes were considered as potential epilepsy-associated genes. The evidence supporting their gene-disease associations varies. Most of these genes had only a single case reported and thus warrant further studies to validate the gene-disease associations. In contrast, clinical and experimental evidence has been obtained supporting the gene-disease association in several genes, such as *UNC13B*. Variants in *UNC13B* have been recurrently identified in unrelated families with partial (focal) epilepsy and/or febrile seizures [26,50]; *Unc13b* knockdown in Drosophila increased seizure-like behavior and firing of neurons in electrophysiological recordings [26]; and other evidence, such as lethality by genetic knockout, high pRec (1.0), expression profile, and vesicle-release related function (https://gdap.org.cn/), also support the gene-disease association.

We listed the top 100 frequently reported/identified epilepsyassociated genes that were ranked by mutation number in the HGMD database and case number in the China Epilepsy Gene 1.0 project, which

Table 3 Epilepsy-related genes (n = 974).

Epilepsy-relat	ed genes $(n = 974)$.
Inheritance	Genes
AD	ABCC8, ACTA2, ACTB, ACTG1, ACVR1, AFF3, AHDC1, AKAP9, AKT2, ALG10B, ANKH, ANKRD11, ANKRD17, APP, ARCN1, ARID1A, ARID1B, ARID2, ASH1L, ASXL1, ASXL3, ATP1A1, ATP2A2, ATP2B1, ATXN10, AUTS2, BCL11B, BICD2, BICRA, BMP4, BRAF, CACNA1D, CACNA1G, CALM2, CALM3, CAMK2B, CAMTA1, CAV3, CCM2, CDC42, CDK13, CELSR1, CERT1, CHD4, CHD8, CHN1, CIC, CLCN6, CLTC, CNOT2, CNOT3, COMT, CTLA4, DHX16, DIP2B, DLG4, DNAJC5, DNMT1, DNMT3A, DRD3, DYNC1H1, DYRK1A, EBF3, EED, EFTUD2, EHMT1, ELN, EZH2, FAM111A, FBX011, FGF8, FGFR2, FGFR3, FXYD2, GARS1, GATA6, GCK, GFAP, GJA1, GLI2, GLI3, GLUD1, GNA11, GNB1, HIVEP2, HMBS, HNF1B, HTR2A, HTRA1, INSR, IRF3, ITPR1, KANSL1,
	KAT6A, KCNA1, KCNE1, KCNE2, KCNH1, KCNH2, KCNJ11, KCNJ5, KCNJ6, KCNK4, KCNQ1, KCNQ5, KDM1A, KDM3B, KDM4B, KIF1A, KMT2A, KMT2D, KMT5B, KRAS, KRIT1, LMX1B, LRRK2, MAF, MAGEL2, MAPZK1, MAPK1, MAPRE2, MAPT, MBD5, MED13, MED13L, MEN13, MN1, MSX2, MYRF, MYT1L, NAA15, NF1, NFE2L2, NF1A, NIPA1, NIPBL, NOD2, NONO, NOTCH1, NOTCH2, NOTCH2NLC, NOTCH3, NPTX1, NR2F1, NSD1, NSD2, OTX2, PACS1, PAK2, PDYN, PHOX2B, PKD1, POLG2, POMP, PPOX, PPP1R12A, PPP2R1A, PPP2R5D, PRKACB, PRNP, PROK2, PSEN1, PSEN2, PTEN, PUF60, PUM1, RAC1, RAD21, RAI1, RANBP2, RBP4, ROBO1, RORA, RRAGD, PROKR2, RNF125, RTN4R, RYR1, RYR2, SATB1, SATB2, SCN5A, SERPINI1, SET, SETBP1, SETD2, SGCE, SHANK3, SIN3A, SLC25A4, SMAD2, SMARCA2, SMARCE1,
	SNORA31, SNTA1, SON, SOX2, SOX4, SPAST, SPECC1L, SPOP, SPTBN1, SPTLC2, SRCAP, SRP54, STAG1, STUB1, SYN2, SYNGAP1, TBK1, TBL1XR1, TBP, TBX1, TCF4, THRB, TLK2, TNPO2, TOP2B, TP53, TRAF3, TRAF7, TRIM8, TRIP12, TTR, UBTF, VPS4A, WDR37,
AR	ZBTB18, ZEB2, ZMYND11, ZNF292 ($n=234$) AASS, ABAT, ABCA2, ABHD12, ABHD16A, ACADM, ACADS, ACADSB, ACO2, ACOX1, ACY1, ADA2, ADAMTSL2, ADD3, ADK, ADPRHL2, ADSL, AFG3L2, AGA, AGL, AGPS, AHI1, ALDH18A1, ALDH3A2,
	ALDH4A1, ALDH5A1, ALDOB, ALG1, ALG11, ALG12, ALG14, ALG2, ALG3, ALG6, ALG8, ALG9, ALKBH8, ALPL, AMACR, AMFR, AMT, ANK3, ANO10, ANTXR1, AP3D1, AP4B1, AP4E1, AP4M1, AP4S1, APOPT1, ARG1, ARHGDIA, ARMC9, ARNT2, ASAH1, ASL, ASNS, ASPA, ASS1, ATAD1, ATG7, ATIC, ATM, ATP13A2, ATP5F1A, ATP5F1D, ATP5F1E, ATP5P0, ATP6V0A2, ATP7B, ATPAF2, AUH, B3GALNT2, B4GAT1, B9D2, BCKDHA, BCL10, BCS1L, BLOC1S6,
	BOLA3, BRCA2, BSCL2, BTD, BUB1B, C12orf57, C2orf69, CAMLG, CARD9, CARS2, CASQ2, CC2D2A, CCBE1, CCDC88A, CD59, CDK10, CEP164, CHKB, CKAP2L, CLDN16, CLN3, CLN5, CLN6, CLN8, CLPP, CNTNAP1, COG2, COG4, COG5, COG6, COG7, COL18A1, COLGALT1, COPB1, COQ2, COQ4, COQ5, COQ6, COQ8A, COQ9, COR01A, COX10, COX11, COX15, COX411, COX6B1, COX8A, COXFA4, CPS1, CPT1A, CRB2, CRBN, CRIPT, CRLF1, CRLS1, CRYAB, CSPP1, CTC1, CTH,
	CTSA, CTSD, CTSF, CYB5R3, CYP27A1, CYP27B1, D2HGDH, DCAF17, DDX59, DHCR24, DHFR, DHTKD1, DLD, DNAJC12, DNAJC21, DNAJC6, DOCK6, DOCK8, DOLK, DPAGT1, DPH1, DPM1, DPM2, DPM3, DPP9, DPYD, DPYS, DSTYK, EARS2, ECHS1, ECM1, EGF, EIF3F, ELOVL4, ELP2, EMC1, EN1, ENTPD1, EPG5, ERCC6, ERLIN2, ESCO2,
	ETFDH, ETHE1, EXTL3, F2, FA2H, FADD, FAR1, FARS2, FARSB, FASTKD2, FAT4, FBXL4, FCHO1, FCSK, FDFT1, FH, FIT2, FKIN, FLVCR2, FMN2, FOLR1, FOXRED1, FTO, FUT8, GALC, GALNT2, GAMT, GBA, GCDH, GCH1, GCSH, GET4, GFER, GFM1, GFM2, GGT1,
	GJC2, GLB1, GLDC, GLE1, GLRB, GLUL, GLYCTK, GM2A, GMPPB, GNB5, GNPAT, GOLGA2, GPAA1, GPSM2, GPX4, GRM1, GRN, GSS, GTPBP3, GUCY1A3, GYS1, GYS2, HACE1, HADHSC, HAX1, HERC2, HEXA, HGSNAT, HHAT, HIBCH, HINT1, HLCS, HMGCL, HMGCS2,
	HMOX1, HPD, HSD17B4, HTRA2, IARS1, IARS2, IBA57, IFNAR1, IFNAR2, IFT140, IL6ST, INPP5K, IQSEC1, IREB2, ISCA1, ISCA2, ISG15, JAM3, KCNJ10, KIAA0753, KIF7, KLHL7, KPTN, L2HGDH, LAMA2, LAMA5, LARS1, LARS2, LCP2, LETM1, LGI3, LGI4, LIAS, LIG3,
	LINGO1, LIPT2, LMAN2L, LMBRD1, LRP2, LRPPRC, LSM11, LSS, MAGI2, MAN1B1, MANBA, MAPKAPK5, MBOAT7, MCCC1, MCCC2, MCM3AP, MCM8, MDM2, MED23, MED25, MEGF10, METTL23, METTL5, MFF, MFSD8, MGAT2, MGP, MICOS13, MICU1, MIPEP,
	MKS1, MLYCD, MMAA, MMACHC, MMADHC, MOCS1, MOCS2, MOGS, MPC1, MPDU1, MPI, MPV17, MRAP, MRM2, MRPL12, MRPS22, MTFMT, MTHFD1, MTHFR, MTO1, MTR, MTX2, MYMK, MYO1H, MYO5A, NADK2, NAGA, NAGLU, NANS, NARS2, NAXD,

NAXE, NBAS, NDST1, NDUFA13, NDUFA2, NDUFA6, NDUFA8,

NDUFAF2, NDUFAF3, NDUFAF4, NDUFAF5, NDUFAF6, NDUFAF8,

NDUFB8, NDUFC2, NDUFS3, NDUFS4, NDUFS7, NDUFS8, NDUFV1,

Table 3 (continued)

Inheritance Gen

NDUFV2, NEU1, NFS1, NGLY1, NIN, NNT, NPC1, NPC2, NRXN1, NSIIN2 NSIIN3 NTRK1 NIIRPI NIIP133 NIIP214 OGDH OPA3 ORAI1, OSGEP, OTUD6B, P4HTM, PAH, PANK2, PARN, PC, PCCA, PCCB, PCDH15, PCK1, PCNT, PCSK1, PCYT2, PDE10A, PDE2A, PDHB, PDHX, PDP1, PDSS2, PET100, PEX1, PEX10, PEX12, PEX13, PEX14, PEX19 PEX2 PEX26 PEX3 PEX5 PEX6 PEX7 PGAP2 PGAP3 PGM1 PGM3, PHGDH, PHKG2, PIBF1, PIGC, PIGF, PIGH, PIGL, PIGM, PIGO, PIGV, PIGW, PIGY, PITRM1, PLA2G6, PLVAP, PMM2, PMPCB, PNPLA8, PNPT1, POLE, POLG, POMC, POMGNT1, POMK, POMT1, POMT2 PPA2 PPCS PPP2R3C PPT1 PRDX1 PRF1 PRKDC PRMT7 PRODH, PRORP, PROS1, PSAP, PSAT1, PSMB8, PSPH, PTRH2, PTS, QDPR, RAB18, RAB27A, RAB3GAP1, RAB3GAP2, RAPSN, RBCK1, RBM8A, RFT1, RMND1, RNASEH2A, RNU12, RNU7-1, ROGDI, RPIA, RRM2B, RSRC1, RTN4IP1, RUBCN, RUSC2, SACS, SARDH, SBF1, SCO2, SCYL1, SCYL2, SDHA, SDHB, SDHD, SELENOI, SERAC1, SGPL1, SGSH, SLC12A1, SLC12A3, SLC17A5, SLC19A2, SLC19A3, SLC25A1, SLC25A10, SLC25A13, SLC25A15, SLC25A19, SLC25A20, SLC25A3, SLC25A36, SLC25A42, SLC26A4, SLC28A1, SLC31A1, SLC33A1, SLC35A1, SLC35A3, SLC35C1, SLC39A8, SLC44A1, SLC45A1, SLC46A1, SLC5A6, SLC6A19, SLC9A1, SMARCAL1, SMC5, SMG9, SMO, SNAP29, SNX14, SOBP, SPTSSA, SQOR, STAR, STAT1, STAT2, STT3A, STT3B, STX11, STXBP2, SUCLA2, SUCLG1, SUMF1, SUOX, SURF1, SYT14, TAF13, TAF2, TANGO2, TARS2, TASP1, TBC1D20, TBCE, TBX19, TCIRG1, TDP1, TDP2, TECPR2, TELO2, TGFB1, THG1L, TIMM50, TIMMDC1, TMCO1, TMEM165, TMEM67, TMEM70, TP53RK, TPK1, TPP1, TRAPPC11, TRAPPC12, TRAPPC9, TRIP13, TRIT1 TRMT1 TRMT5 TRNT1 TRPM6 TSFM TSPYI.1 TWNK TXN2, TYROBP, UCHL1, UGT1A1, UNC13D, UPB1, UQCC2, USP18, USP53, VARS2, VPS13A, VPS13B, VPS13D, VPS35L, VPS41, WASHC4, WFS1, WIPI2, XPNPEP3, YARS1, YRDC, ZBTB11, ZFYVE26, ZMPSTE24, ZNFX1 (n = 605)AD, AR APOE, CAMK2A, CFH, CLPB, CNNM2, CPT2, DNM1L, FTL, GCM2, GLRA1, GRIA1, HESX1, HTT, IFIH1, OPLAH, POLRMT, PROC, PTH, RNF213, SLC16A1, SPR, TET3, TICAM1, TLR3, TREX1, ZFP57 (n = 26) AMER1, ATRX, BCOR, CLCN4, FMR1, GDI1, HCCS, HNRNPH2, XLD HSD17B10, IQSEC2, KDM6A, MSL3, NDUFB11, NEXMIF, NHS, PDHA1, USP9X, WDR45 (n = 17)XLR ABCD1, AFF2, AIFM1, ANOS1, AP1S2, ATP6AP1, ATP6AP2, ATP7A, AVPR2, BCAP31, BRWD3, CHRDL1, CLIC2, CUL4B, DKC1, DLG3, EBP, EIF2S3, FAM50A, FGD1, FTSJ1, GK, GRIA3, HCFC1, HPRT1, IDS. IKBKG, IL1RAPL1, KDM5C, KIF4A, KLHL15, LAGE3, MAOA, MBTPS2, MECP2, MED12, MID2, NDP, NDUFA1, OCRL, OFD1, OTUD5, PAK3, PGK1, PHF6, PHF8, PQBP1, RAB39B, RBM10, RPL10, SLC6A8, SMS, SSR4, STS, SYP, TAF1, THOC2, TMLHE, UBE2A, UPE3B (n = 60)XL CNKSR2, DDX3X, FRMPD4, GLRA2, NLGN3, OTC, RNF113A, SLC9A6, TFE3, XK, ZDHHC9 (n = 11)DR KNSTRN+PIK3CD (n=2)ACSF3. BCKDK. CACNG2. CEP290. COG8. FGFR1. GNAO. GPHN. IIN HADHA, HADHB, IDH2, KCNK9, MTSS2, NAT8L, NRAS, PIK3CA,

AD, autosomal dominant; AR, autosomal recessive; DR, Digenic recessive; UN, unknown; XL, X-linked; XLD, X-linked dominant; XLR, X-linked recessive.

SLC1A3, SLC9A9, ZFHX3 (n = 19)

may be referred in clinical practice. While severe epilepsies such as epileptic encephalopathy are more commonly targeted in clinical and genetic studies, attention is advocated for genes associated with common epilepsies, which is potentially significant in the clinical management of patients, e.g., *UNC13B* [26] and *FAT1* [35]. The patient with the biallelic *UNC13B* variant presented with frequent daily seizures with abnormalities of the hippocampus but achieved seizure-free status without surgery [26]. The patients with *FAT1* variants showed relapse of epilepsy after long-term seizure-free status, which could potentially be explained by the genetic expression (dependent) stage of *FAT1* [35], offering an example of how to consider the optimal duration of therapies.

Regarding whether a gene is associated with a phenotype, two frameworks have been developed previously by Clinical Genome Resource (ClinGen) [51] and Ambry Genetics [52]. The two frameworks score publicly available clinical genetic and experimental evidence, and the scoring on clinical genetic evidence was generally based on the number of cases/variants or statistical data. ClinGen has curated 2265

Table 4 Potential epilepsy-associated genes (n = 1440).

Source HGMD Gene

AACS, AARS2, AARSD1, AATK, ABCA10, ABCA5, ABCA6, ABCA7, ABCB1, ABCB11, ABCB9, ABCC12, ABCC2, ABCC4, ABCC6, ABCG8, ACADVL, ACAP3, ACCS, ACD, ACER2, ACMSD, ACOT7, ACP4, ACSM2A, ACSM5, ACTA1, ACTC1, ACY3, ADA, ADAM21, ADAMTS16, ADAMTS17, ADAMTS4, ADAR, ADCY1, ADCY7, ADCYAP1R1, ADGRE2, ADGRF4, ADPRM, ADRA1B, ADRA2B, AFAP1L1, AFAP1L2, AFF4, AGAP1, AGBL4, AGFG2, AGTR2, AKAP6, AKAP7, ALG10, ALK, ALKBH2, ALPG, ALX4, AMBRA1, AMHR2, AMZ1, ANK1, ANK2, ANKRD16, ANKRD30A ANKRD33B, ANKRD36, ANKRD54, ANO3, ANO7, ANXA6, AOC1, AP1B1, AP5B1, AP5Z1, APBA2, APBA3, APC, APOB, APOC2, APRT, ARAP3, AREL1, ARF3, ARGLU1, ARHGAP21, ARHGAP24, ARHGAP32, ARHGAP35, ARHGEF10, ARHGEF15, ARHGEF18, ARHGEF7, ARNTL2, ARRB2, ARSH, ASB11, ASGR2, ASPH, ASTE1, ASTL, ASTN1, ASTN2, ATAD3B, ATAD3C, ATAT1, ATG12, ATG16L2, ATG2A, ATG3, ATP13A5, ATP2A1, ATP2B3, ATP6V0D1, ATP6V1B2, ATP8A2, ATP8B2, ATPAF1, ATRN. ATRNL1, AZI2, BAG2, BAIAP2, BAZ1A, BBS2, BCAM, BCL11A, BCL2L12, BCLAF1, BCR, BDNF, BEAN1, BEST2, BEX3, BHLHE40, BHLHE41, BIRC6, BMP10, BMP5, BMP6, BRD7, BSN, C11orf65, C1orf54, C1orf87, C1QTNF12, C1QTNF7, C20orf194, C2CD3, C3, C3orf20, C7orf50, C9orf72, CABP4, CADM3, CADPS2, CALN1, CAMK2G, CAMTA2, CAND1, CAPN1, CAPN10, CAPN11, CAPN3, CAPZA1, CAPZA2, CARD10, CARD14, CARNS1, CASP8AP2, CAVIN2, CBL, CC2D1A, CCDC121, CCDC127, CCDC141, CCDC15, CCDC17, CCDC178, CCDC186, CCDC96, CCKBR, CCN3, CD177, CD93, CDAN1, CDC123, CDC25C, CDC42EP1, CDC7, CDH11, CDH12, CDH13, CDH15, CDH22, CDH4, CDH6, CDH8, CDK20, CDK5R2, CDK5RAP1, CELF4, CELSR3, CENPQ, CEP120, CEP170, CEP170B, CEP68, CEP76, CEP89, CETP, CFAP69, CFAP91, CHADL, CHCHD1, CHCHD6, CHD11, CHD7, CHGA, CHL1, CHMP4C, CHRFAM7A CHRM1, CHRNA3, CHRNA7, CHRNB3, CHST2, CINP, CLCN1, CLDN23, CLIP1, CLOCK, CLRN3, CLSTN1, CLTCL1, CMYA5, CNBP, CNIH3, CNIH4, CNNM3, CNNM4, CNPY2, CNTD1, CNTN1, CNTN5, CNTN6, COASY, COL10A1, COL11A2, COL12A1, COL13A1, COL17A1, COL20A1, COL21A1, COL2A1, COL5A3, COL6A2, COL6A3, COL7A1, COL9A2, COLQ, COPB2, CORO7, COX6A1, CP, CPEB2, CPNE2, CPNE6, CPQ, CRH, CRHR1, CRIM1, CSAD, CSMD1, CSMD3, CSNK1E, CSNK1G1, CTBP2, CTDP1, CTNNA3, CTNNB1, CTNND2, CTR9, CTTNBP2, CUBN, CUX1, CWF19L1, CXCR4, CYB561A3, CYFIP1, CYP2A13, CYP51A1, CYREN, DAB2IP, DAG1, DAGLA, DAGLB, DAP3, DCAF13, DCAF15, DCC, DCHS1, DCLK2, DDX31, DDX43, DDX49, DDX60, DDX60L, DENND1A, DENND2B, DENND2C, DENR, DGAT1, DGKD, DGKE, DGKG, DGKZ, DHODH, DHRS3, DHRS9, DIAPH3, DIP2C, DLAT, DLG1, DLG2, DLGAP2, DLGAP4, DMBX1, DMD, DMXL1, DNA2, DNAH1, DNAH11, DNAH14, DNAH6, DNAH7, DNAJA1, DNAJB8, DNAJC15, DNM3, DOCK10, DOK7, DPP8, DRP2, DSCAM, DSP, DTWD2, DTX3, DUOX1, DUOX2, DUS1L, DUS2, DUS3L, DVL2, DYNC111, DYNC2H1, DYNC2I1, DYSF, DZIP1, ECRG4, ECT2, EDEM2, EDN3, EEF1AKMT3, EFCAB1, EFNA3, EFNB2, EIF2AK3, EIF3E, ELFN1, ELMO1, ELOA, EML4, EML5, EML6, ENPP7, ENTPD3, EP300, EP400, EPB41L3, EPHA2, EPHA5, EPHB1, EPHB2, EPHB6, ERAP2, ERBB4, ERMAP, ESF1, ESPNI, ESRP1, ESYT2, ETV1, ETV2, EVPI, EXD3, EXOC6. EXOC6B, EYA1, EYA3, F13A1, F5, FAAH2, FAAP20, FAHD2A, FAM114A1, FAM131C, FAM149A, FAM151A, FAM170A, FAM180B, FAM186A, FAM20C, FAM219A, FAM71B, FAM89A, FANCA, FANCC, FASN, FAT1, FAT2, FAT3, FBN2, FBXO22, FBXO25, FBXO34, FBXO40, FERMT3, FFAR1, FFAR2, FGB, FGD6, FGF1, FGGY, FHIT, FHL1, FKBP11, FLNC, FMC1, FN1, FN3KRP, FNDC7, FNTB, FOSL1, FOXA2, FOXRED2, FPR2, FRMD1, FRMPD2, FSIP2, FSTL5, FURIN, FUT1, FYTTD1, FZD9, GABBR1, GABPB2, GABRA4, GABRA6, GABRE, GABRG1, GABRG3, GABRP, GABRO, GABRR2, GADD45GIP1, GALNS, GALNT1, GANAB, GAPDHS, GAPT, GARNL3, GAS2L2, GATA3, GATA4, GATM, GBE1, GDAP1, GEMIN2, GFRA1, GHRL, GIPC1, GJD2, GLG1, GLP1R, GMPPA, GNA12, GNPTAB, GOLGA7B, GORAB, GP5, GPI, GPM6A, GPR179, GPR37L1, GRAMD1C, GRAMD4, GRB10, GREB1L, GRHL2, GRIK1, GRIK4, GRIP1, GRM3, GRM4, GRM8, GSPT2, GSTCD, GSTM1, GTF2A2, GTF2IRD2, GTF3C3, GTF3C5, GUCY2D, GUSB, HACD3, HAP1, HARS1, HDAC1, HDAC3, HDAC8, HDAC9, HEATR5B, HECW1, HELZ, HEPHL1, HES1, HES4, HEXB, HGD, HIP1, HIRA, HIVEP1, HMCN1, HMGB4, HNRNPA1, HNRNPAB, HOXA6, HPS3, HR, HS2ST1, HSPA12A, HSPA1L, HSPE1, HTR1A, HTR1F, HTR2C, HTR6, HYDIN, ICAM3, ID2, IDH1, IDUA, IFNA8, IGFALS, IGHMBP2, IGSF6, IGSF8, IL10, IL12RB1, IL1B, IL1RN, IL27RA, II.6. IMMP21. IMPDH2. INA. INCENP. ING5. INO80. INPP4A. IP6K3. IOCC. IQGAP2, IQGAP3, ITFG2, ITGA4, ITGA5, ITGB1BP1, ITIH4, ITLN1, ITPRID2, JKAMP, JMJD1C, JRK, JUP, KAT14, KAT6B, KATNAL2, KATNIP, KBTBD13, KCNAB1, KCNAB2, KCNC3, KCND1, KCND2,

KCND3, KCNG1, KCNH5, KCNIP1, KCNJ2, KCNMB1, KCNMB3, KCNN3,

Table 4 (continued)

Genes

Source

KCNV2, KCTD13, KCTD14, KCTD19, KCTD3, KDM3A, KDM5A, KDM5B, KDR KEL KERA KHK KIAA0100 KIAA0319L KIE13B KIE1C KIE3C KIF5B, KIF6, KIRREL3, KLC2, KLHDC4, KLHDC9, KLHL17, KLHL20, KLHL41, KLHL42, KLRG2, KMT2B, KNL1, KNTC1, KPNA7, KRT1, KRT4, KRT6B, LAMA1, LAMB2, LAMB3, LAMC2, LARGE1, LARP1B, LARP4, LARP7 LCTL LDLR LEAP2 LEPR LGALS3RP LIFR LINS1 LIPN LIPT1 LMAN1L, LMNTD2, LNX1, LORICRIN, LOXL3, LPA, LPAR2, LRFN2, LRFN5, LRIG1, LRP12, LRP1B, LRRC24, LRRC37A3, LRRC4, LRRC40, LRRC4C, LRRC74A, LRRD1, LRRIQ4, LSM1, LYPLA2, LYRM1, LZTR1, MAD11.1 MAGED1 MAGI2 MAGI3 MAP2 MAPK10 MAPK3 MARCHF9, MAST4, MASTL, MATN2, MATN4, MBD3L1, MBL2, MBOAT2, MC3R, MC4R, MCF2L, MCM4, MCMDC2, MCOLN1, MCRS1, MDN1, ME3, MED30, MEGF6, MEIS2, MEPCE, METRN, METTL21A, METTL22, MFN2, MFSD6, MGA, MICAL1, MICAL3, MICALL1, MILR1, MINDY2, MKI67, MKRN2. MLLT1. MLLT3. MMP21. MMP23B. MMP27. MMRN2. MMUT. MOCOS, MORC2, MPL, MPLKIP, MPP4, MPP7, MR1, MRI1, MRPL35, MRPL38, MRPL54, MRPS27, MS4A10, MSC, MSH6, MSR1, MT1A, MTMR9, MUC16, MUC4, MUSK, MUTYH, MYH11, MYH8, MYO16, MYO1B, MYO3B, MYO5C, MYO6, MYOCD, MYOF, MYOM2, MYPN, MYSM1, MYT1, MZT2B, N4BP1, NBEAL1, NCAPD2, NCBP3, NCEH1, NCKAP5, NCOR2, NDC80, NDUFAF1, NDUFB3, NDUFB7, NEB, NECAB1, NECTIN2, NECTIN4, NEDD4, NEIL1, NELL1, NELL2, NEMF, NF2, NFIB, NFIC. NFKBIZ. NFU1. NID1. NID2. NINL. NIPA2. NKAIN3. NLGN1. NLGN4X, NLRP8, NMD3, NMT1, NODAL, NOL3, NOLC1, NOMO2, NOS3, NOTUM, NPAS2, NPAT, NPC1L1, NPR1, NPR2, NQO1, NRG2, NRG3, NRXN2, NRXN3, NTAN1, NTF3, NTN3, NTNG1, NUP160, NUP85, NUTM2F, NUTM2G, NXNL1, NXPE4, OAS2, OCA2, OCSTAMP, OGT. OLAH, OLFM1, OLFML2A, OLFML3, OMG, OPA1, OPCML, OPN1SW, OPRM1, OR10G9, OR10H2, OR10Q1, OR10Z1, OR11A1, OR4N4, OR51A7, OR51D1, OR5A1, OR5L2, ORC4, ORMDL1, OSBP2, OSBPL5, OSTM1 OTOP1 OTUD7A OVCH2 P2RX7 PAK6 PAPPA PAPPA2 PAQR7, PARP10, PAX3, PAX5, PAX6, PCDH11X, PCDH7, PCDH9, PCDHB4, PCDHGA4, PCDHGA5, PCDHGA8, PCED1A, PCGF1, PCIF1, PCNX2, PCNX4, PCSK7, PDCD1, PDE6B, PDIA6, PEG3, PER1, PFAS, PFKFB1, PGAP6, PGBD2, PGBD3, PGK2, PI15, PI4K2A, PIK3AP1, PIK3R4, PIP5K1C, PIR, PITHD1, PKDREJ, PKHD1L1, PLA2G4A, PLAC8L1, PLB1, PLCB4, PLCG2, PLCL1, PLD5, PLEC, PLEKHG6, PLEKHH1, PLIN3, PLIN5, PLK1, PLXDC2, PLXNA3, PLXNB1, PML, PMM1, PMP22, PMPCA, PNLIP, PNPLA6, PODXL, POGLUT2, POLI, PON2, PORCN, PPARGC1B, PPFIA3, PPM1D, PPP1CB, PPP1R12C, PPP1R3C, PPP2R3A, PPP5C, PPP6R2, PRAG1, PRAMEF1, PRCP, PREB, PREX2, PRICKLE2, PRIMA1, PRKCG, PRKD1, PRKN, PRL, PRMT5, PROCA1, PROM1, PRPF8, PRR22, PRRC2C, PRTG, PSMD7, PTCHD1, PTER, PTGER2, PTGS2, PTH1R, PTK2, PTPN11, PTPRD, PTPRE, PTPRM, PTPRZ1, PYCR1, R3HCC1L, RAB11A, RAB2B, RAB3D, RAB40C, RAD23A, RAD52, RAD54L, RAG2, RANBP3L RANGAP1, RAP1GAP2, RAPGEFL1, RASA1, RASEF, RBFOX1, RBFOX3, RBM34, RBPJ, RBSN, RCL1, RDH12, RDH5, RDX, RECQL5, RFX3, RFX4, RGL3 RHRDF1 RICTOR RIMBP2 RIN2 RIOK2 RIPOR1 RLBP1 RNASEH2C, RNF34, ROBO4, RPAP1, RPGRIP1L, RPH3A, RPL4, RPS6KB2, RRM1, RRNAD1, RRP1B, RRP9, RSRC2, RTN2, RTP1, RUFY1, RUNDC3A, RXFP1, RXFP4, RYR3, S1PR4, SAMD4B, SAMD7, SAMHD1, SATL1, SBSPON, SCAF4, SCAMP5, SCN10A, SCN4A, SCN7A, SCN9A, SCNM1, SCNN1B, SCNN1D, SDF2, SDK1, SEC23A, SEC24B, SEC24D, SEMA3E, SEMA5A, SEPTIN8, SERHL2, SETD5, SETDB1, SETX, SEZ6, SF3B3, SFR1, SGCG, SH3BGR, SH3GLB1, SH3RF1, SHANK1, SHANK2, SHROOM4, SIDT1, SIRT3, SKI, SLC12A2, SLC12A7, SLC15A3, SLC16A12, SLC16A2, SLC17A2, SLC17A3, SLC18A2, SLC1A1, SLC1A5, SLC1A7, SLC22A1, SLC22A20P, SLC22A5, SLC23A3, SLC24A4, SLC25A39, SLC25A40, SLC26A1, SLC26A2, SLC2A12, SLC30A3, SLC30A8, SLC32A1, SLC35E4, SLC37A2, SLC37A3, SLC38A10, SLC39A13, SLC3A1, SLC3A2, SLC44A2, SLC4A10, SLC4A3, SLC5A12, SLC5A8, SLC6A3, SLC6A4, SLC6A9, SLC7A11, SLC7A13, SLC7A3, SLC7A4, SLC9A4, SLCO4C1, SLTM, SMARCA1, SMCHD1, SMCO2, SMG6, SMOC2, SMPDL3A, SNAP23, SNAP25, SNAP47, SNAP91, SNRNP40, SNX19, SNX27, SOAT1, SOCS2, SOHLH1, SORCS3, SOS1, SOWAHB, SOWAHC, SOX10, SOX11, SP4, SPAG4, SPAG9, SPANXA2, SPATA16, SPATA7, SPATA8, SPDL1, SPG11, SPHKAP, SPIRE2, SPRED2, SPRY4, SPTB, SQLE, SQSTM1, SRD5A3, SREK1, SRGAP2, SSPOP, SSTR1, SSTR5, ST6GALNAC3, ST7, ST8SIA2, STAB1, STAB2, STAG2, STARD13, STARD9, STC1, STK11, STK31, STRAP, STRIP1, STRIP2, STXBP5L, STYXL1, SUCLG2, SUCO, SULT6B1, SUN1, SV2A, SVIL, SWSAP1, SYCP2, SYN3, SYNCRIP, SYNE1, SYNE2, SYNJ2, SYNM, SYT2, SYT4, SYT9, TACC2, TACR2, TAF1L, TAMALIN, TAP1, TARS1, TBC1D10B, TBC1D8, TBC1D8B, TBL2, TBX18, TCHHL1, TCOF1, TDO2, TEX33, TEX45, TFAM, TFAP2E, TGFB111, TGIF1, TH, THAP11, THEMIS, THOP1, THUMPD3, TIGD4, TIMELESS, TK2, TKT, TMC7, TMED7, TMEM132B, TMEM139, TMEM229B, TMEM61, TMEM63A,

(continued on next page)

China Epilepsy Gene 1.0 project (case

Table 4 (continued)

Source Genes

TMEM63B, TMEM68, TMPRSS6, TMPRSS9, TNC, TNIP2, TNIP3, TNK2, TNN13K, TNR, TNS1, TNS2, TOM1L2, TOP1MT, TOP3B, TPTE2, TRAZB, TRAF5, TRANK1, TRIM24, TRIM29, TRIM50, TRIM66, TRIM69, TRIP11, TRIP4, TRMT44, TRPC6, TRPV5, TSG101, TSNARE1, TSPAN5, TSPAN7, TSSC4, TTBK2, TTC21B, TTC28, TTC3, TTC30A, TTC39A, TT11, TUBA3E, TUBA8, TULP2, TUT1, TWSG1, TXNRD1, TYK2, U2AF2, UBR4, UBR5, UGGT2, UGT1A9, UGT2B4, UHRF1BP1, UIMC1, UNC13A, UNC13B, UPF1, UPP2, UQCRFS1, USH1C, USH2A, USP19, USP2, USP24, USP28, USP29, USP34, USP36, USP8, UTP20, VAMP5, VCL, VIL1, VOPP1, VPS13C, VPS18, VPS35, VPS39, VPS45, VRK2, VWA3B, VWA8, WDFY3, WDHD1, WDR3, WDR41, WDR59, WDR72, WDR75, WDR91, WHAMM, WNT4, WRAP53, WRN, XDH, XRCC1, YAP1, YTHDC2, YWHAE, YY1, YY1AP1, ZADH2, ZBTB20, ZCCHC8, ZDHHC1, ZDHHC15, ZFR2, ZGPAT, ZGRF1, ZHX3, ZKSCAN2, ZNF105, ZMF208, ZMF221, ZNF248, ZNF302,

ZNF317, ZNF318, ZNF343, ZNF345, ZNF354A, ZNF385A, ZNF397,

ZNF408, ZNF420, ZNF433, ZNF44, ZNF443, ZNF517, ZNF527, ZNF536,

ZNF565, ZNF570, ZNF585B, ZNF598, ZNF654, ZNF655, ZNF675, ZNF727,

ZNF774, ZNF789, ZNF813, ZNF814, ZNF844, ZPBP2, ZYG11A, ZYX (n =

PubMed

1367)

AAAS, ACAD9, ADORA2A, AIPL1, ARL13B, ARL6IP1, ARSB, ATP6, BRD2, CD9912, CELSR2, CLCN7, EEF1B2, ELMO2, ENG, FAN1, FBP1, FBXO7, FCN3, GJB2, GLA, GRINA, GSN, HIKESHI, HOXD, IDH3A, KIAA0586, KLF13, LONP1, MATR3, MBTPS1, ME2, MEFV, MTCO2, MTTF, MTTK, MTTL1, MTTM, MYH1, MYH9, MY018A, OSTC, PAX2, PAX7, PCDHG, PKP2, POLR1A, PRKACA, PRKAG2, PTPN4, RB1, RHEB, SCA2, SHOC2, SLC16A4, SLC16A7, SLC25A2, SLC29A1, SLC6A12, TAF1C, TAT, TBC1D22A, TECTA, TENM1, TOR1A, TRAPPC2L, TRPM1, TUBGCP5, TUSC3, UBA1, UBP1, UNC79, ZNF385B (n = 73)

Bold italics, with a number of mutations identified in the epilepsy phenotypes >5.

genes on their gene-disease associations, including 95 genes in the expert panel list of epilepsy (https://search.clinicalgenome.org/kb/gene-validity, until August 2023), while Ambry Genetics curated the gene-disease relationships on 22 genes [52].

We have developed a pathogenic potential and pathogenic feature assessment (PPA) system to evaluate whether (pathogenic potential) and how (pathogenic feature) a gene is associated with a phenotype with criteria on clinical evidence that include phenotype (spectrum) specificity, inheritance pattern, pathogenic genotype, and genotypephenotype correlation, and on experimental evidence including gene expression profile, and gene KO consequence, and other experiments directly related to gene-disease association [53]. Using the PPA framework, 875 epilepsy-associated genes were assessed with available clinical data (until April 2022), of which 617 genes were evaluated as "pathogenic", 150 genes were evaluated as "possible pathogenic", and 108 genes were "to be confirmed" in pathogenicity. The details of evidence in evaluating the gene-disease association are presented in the Genetic Dependence and Pathogenicity (GD&P) database (https://www. gdap.org.cn/statistic/phenotype). The pathogenic potential and pathogenic features of each gene are essential parts of the gene-disease association and fundamental bases for evaluating the pathogenicity of

Recently, two genetic databases for epilepsy have been established. The SAGAS database refers to genes associated with monogenic and polygenic seizures/epilepsy in multiple species [5], among which almost 1000 genes were not included in the present epilepsy-associated gene list due to insufficient clinical evidence. However, attention is required to these genes in searching for potential novel epilepsy genes in further studies. The Gene4epilepsy database is based on previously used epilepsy gene panels and two research resources. Almost 90% of the genes are related to developmental and/or epileptic encephalopathy [6], to which attention is required in application for identifying causative genes in patients with common epilepsies.

This study has several limitations. Generally, the gene-specific burden of variants should be derived from the result of an epidemiological investigation. The data from the HGMD database and the China

Table 5

Top 100 genes reported to be associated with epilepsy in HGMD Database and China Epilepsy Gene 1.0 project.

number)

HGMD Database (mutation number) SCN1A (3374), MECP2 (939), KCNQ2 (731), SCN2A (662), CDKL5 (523), PCDH19 (522), STXBP1 (404), SCN8A (394), DEPDC5 (298), ALDH7A1 (261), SYNGAP1 (231), GRIN2A (222), TPP1 (209), KCNT1 (180), PRRT2 (180), CHD2 (176), CLN6 (174), SLC2A1 (154), FOXG1 (152), TBC1D24 (137), CACNA1A (134), EPM2A (121), PPT1 (121), GABRG2 (119), PIGN (119), GABRB3 (112), TSC2 (110), SLC6A1 (96), WWOX (96), KCNB1 (95). NHLRC1 (92), POLG (89), ARX (88), KCTD7 (86), <u>IQSEC2</u> (83), <u>HNRNPU</u> (81), SCN9A (81), CLN3 (80), GABRA1 (80), GRIN2B (77), NEXMIF (77), CACNA1H (76), SLC13A5 (76), CLN5 (74), LGI1 (74), SPTAN1 (71), SZT2 (70), GNAO1 (67), ADGRV1 (66), KCNA2 (66), NPRL3 (65), GABRB2 (62), TSC1 (60), BRAT1 (56), CLN8 (56), KCNQ3 (56), HCN1 (55), NRXN1 (55), PIGA (55), SPATA5 (53), WDR45 (52), CNTNAP2 (50), PNPO (50), SMC1A (50), MEF2C (47), SCN1B (47), EFHC1 (46), SCN3A (46), UBE3A (45), ATP1A3 (44), DNM1 (43), MFSD8 (42), STX1B (41), ARHGEF9 (38), PNKP (38) RARS2 (38) CSNK2B (37) EEF1A2 (37), GRIN1 (36), ZEB2 (36), ATP1A2 (35), MBD5 (34), ITPA (32), NBEA (32), PURA (32), COL4A1 (31), DYRK1A (31), KARS (30), PLPBP (29), RELN (29), WDR26 (29), ADSL (28), CLCN2 (28), PIGT (28), AARS1 (27), ALG13 (27), KCNJ10 (27), SYN1 (27),

SCN1A (58), FLNA (18), PRRT2 (16), TSC1 (13), CELSR2 (11), CHD2 (11), HCFC1 (11), TSC2 (11), ADGRV1 (10), CACNA1A (10), SLC2A1 (10), CACNA1H (9), SCN2A (9), ZFHX3 (9), BCOR (8), GABRG2 (8), KMT2D (8), NF1 (8), PKD1 (8), RELN (8), AFF2 (7), MECP2 (7) MED12 (7) NEXMIF (7) PCDH15 (7), PCDH19 (7), SETD1B (7), ATP7A (6), BSN (6), CDKL5 (6), CELSR1 (6), DEPDC5 (6), FAT1 (6), KCNO2 (6), KCNT1 (6), LAMA5 (6), PCNT (6), <u>SCN8A</u> (6), <u>SZT2</u> (6), **TENM1** (6), AIFM1 (5), ALG13 (5), BRWD3 (5), CELSR3 (5), DMXL2 (5), FRMPD4 (5), GRIN2A (5), KCNA1 (5), RYR2 (5), SHROOM4 (5), SYNGAP1 (5), UNC13B (5), ABCA2 (4), APC2 (4), ARX (4), CHD4 (4), CNKSR2 (4), DCHS1 (4), DCX (4), FGD1 (4), GABRA1 (4), IQSEC2 (4), KCNMA1 (4), LRPPRC (4), MGA (4), MID2 (4), NHS (4), NPRL3 (4), PGM3

(4), PI4KA (4), PIGA (4), RYR1 (4),

TBC1D24 (4), THOC2 (4), WDFY3 (4),

ATP1A2 (3), CPA6 (3), DYNC1H1 (3),

GRIN2B (3), HNRNPU (3), IRF2BPL (3),

KDM6A (3), MBD5 (3), MPDZ (3), SBF1

(3), <u>SCN9A</u> (3), <u>SMC1A</u> (3), <u>SYN1</u> (3), TUBGCP6 (3), UNC80 (3), <u>WWOX</u> (3),

YWHAG (3), ATP6VOC (2), UNC79 (2)

EFHC1 (3), GABRB3 (3), GRIA3 (3),

KCNK4 (3), KCNV2 (3), KDM5C (3),

 $^{\#}$ The data were obtained from trio-based whole-exome sequencing analysis on 2757 unrelated cases with epilepsy.

Bold italics, novel epilepsy-associated genes discovered by China Epilepsy Gene 1.0 project.

Underline, identical genes in two databases.

CNKSR2 (26), TANC2 (26)

Epilepsy Gene 1.0 project were just two examples for reference, and attention is required to possible biases in such data. The present gene list did not consider the role of epigenetics and polygenic factors in epilepsy. The 2946 listed epilepsy-associated genes vary in their association with epilepsy, to which attention is required in application.

In conclusion, the present study updates the list of epilepsy-associated genes based on currently available clinical-genetic evidence, which is helpful for gene screening in clinical practice. Recognition of epilepsy-associated genes and their relationships with other abnormal phenotypes is potentially useful for the precise clinical molecular diagnosis and management of patients with epilepsy.

Ethical publication statement

All procedures performed were in accordance with the ethical standards of the institutional committee. The present study was approved by the Ethics Committee of the Second Affiliated Hospital of Guangzhou Medical University.

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Declaration of Competing Interest

The authors have stated that they had no interests that might be perceived as posing a conflict or bias.

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Supplementary materials

Supplementary material associated with this article can be found, in the online version, at doi:10.1016/j.seizure.2023.09.021.

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