

# Channelopathies That Lead to Sudden Cardiac Death: Clinical and Genetic Aspects



Jonathan R. Skinner, MBChB, DCH, MD <sup>a,b,c,\*</sup>,  
Annika Winbo, MD, PhD <sup>a,b,d</sup>, Dominic Abrams, MBChB, MD <sup>e,f</sup>,  
Jitendra Vohra, MD <sup>g,h</sup>, Arthur A. Wilde, MD, PhD <sup>i,j</sup>

<sup>a</sup>The Cardiac Inherited Disease Group, Auckland, New Zealand

<sup>b</sup>Greenlane Paediatric and Congenital Cardiac Services, Starship Children's Hospital, Auckland, New Zealand

<sup>c</sup>Department of Paediatrics, Child and Youth Health, University of Auckland, New Zealand

<sup>d</sup>Department of Neurophysiology, University of Auckland, New Zealand

<sup>e</sup>Inherited Cardiac Arrhythmia Program, Department of Cardiology, Boston Children's Hospital, Boston, MA, USA

<sup>f</sup>Harvard Medical School, Boston, MA, USA

<sup>g</sup>Cardiology Department and Department Of Genomics, The Royal Melbourne Hospital, Melbourne, Vic, Australia

<sup>h</sup>University Of Melbourne, Melbourne, Vic, Australia

<sup>i</sup>Heart Centre AMC, Department of Clinical and Experimental Cardiology, Academic Medical Center, University of Amsterdam, Amsterdam, The Netherlands

<sup>j</sup>Department of Medicine, Columbia University Irving Medical Center, New York, NY, USA

Forty per cent (40%) of sudden unexpected natural deaths in people under 35 years of age are associated with a negative autopsy, and the cardiac ion channelopathies are the prime suspects in such cases. Long QT syndrome (LQTS), Brugada syndrome (BrS) and catecholaminergic polymorphic ventricular tachycardia (CPVT) are the most commonly identified with genetic testing. The cellular action potential driving the heart cycle is shaped by a specific series of depolarising and repolarising ion currents mediated by ion channels. Alterations in any of these currents, and in the availability of intracellular free calcium, leaves the myocardium vulnerable to polymorphic ventricular tachycardia or ventricular fibrillation. Each channelopathy has its own electrocardiogram (ECG) signature, typical mode of presentation, and most commonly related gene. Long QT type 1 (gene, *KCNQ1*) and CPVT (gene, *RyR2*) typically present with cardiac events (ie syncope or cardiac arrest) during or immediately after exercise in young males; long QT type 2 (gene, *KCNH2*) after startle or during the night in adult females—particularly early post-partum, and long QT type 3 and Brugada syndrome (gene, *SCN5A*) during the night in young adult males. They are commonly misdiagnosed as seizure disorders. Fever-triggered cardiac events should also raise the suspicion of BrS. This review summarises genetics, cellular mechanisms, risk stratification and treatments. Beta blockers are the mainstay of treatment for long QT syndrome and CPVT, and flecainide is remarkably effective in CPVT. Brugada syndrome is genetically a more complex disease than the others, and risk stratification and management is more difficult.

## Keywords

Sudden death • Channelopathy • Long QT syndrome • Brugada syndrome • CPVT • Genetics

## Introduction

Forty per cent (40%) of sudden unexpected natural deaths in people under 35 years of age are associated with a negative autopsy and the cardiac ion channelopathies are the prime

suspects in such cases [1]. Often called “SADS” or sudden arrhythmic death syndrome, Long QT syndrome (LQTS), Brugada syndrome (BrS) and CPVT (catecholaminergic polymorphic ventricular tachycardia), are the most commonly found [2–7].

\*Corresponding author at: Green Lane Paediatric and Congenital Cardiac Services, Starship Children's Hospital, Private Bag 92024, Auckland 1142, New Zealand. Tel.: +64 9 3074949 Fax: +64 9 6310785., Email: [j Skinner@adhb.govt.nz](mailto:j Skinner@adhb.govt.nz)

Crown Copyright © 2018 Published by Elsevier B.V. on behalf of Australian and New Zealand Society of Cardiac and Thoracic Surgeons (ANZSCTS) and the Cardiac Society of Australia and New Zealand (CSANZ). All rights reserved.

More than two-thirds of all young sudden cardiac deaths occur at rest or sleep [8,9]; however, a positive genetic result for an ion channelopathy is more likely with a history of exercise or extreme emotion [7]. Among college athletes, SADS is up to three times more common than hypertrophic cardiomyopathy except in African-Americans [10].

Genetic mutations linked to cerebral ion channelopathies causing epilepsy occur in up to 6% of SADS cases [1] and whether these are cardiac or neurologic deaths is not clear, but ion channel dysfunction can present in the brain and heart in the same patient [11,12].

## Presentation Other Than Sudden Death or Cardiac Arrest

Each ion channelopathy has its own electrocardiogram (ECG) signature, and typical mode of presentation (see Figures 1 and 2). They are commonly misdiagnosed as seizure disorders [13,14]. Their coincidental presence may make arrhythmic death more likely in the event of myocardial infarction or illnesses with metabolic disturbance or polypharmacy [15]. Fever-triggered cardiac events should raise the suspicion of BrS [16].

## Prevalence

Long QT syndrome occurs in 1 in 2,000 people with a slight predominance of females, [17] BrS occurs in about 1 in 10,000 people (higher in East Asia) [18], with over 70% being male [19], and CPVT in occurs in approximately 1 in 10,000. Short QT syndrome is very rare indeed and is not reviewed further here. Early repolarisation is an ECG feature of about 5% of the population, but it is suggested that 30% of cases of so-called idiopathic ventricular fibrillation (VF) are due to early repolarisation syndrome (ERS).

## Mechanism of Arrhythmogenesis

### The Cardiac Action Potential and Cellular Excitability

The cellular action potential driving the heart cycle is shaped by a specific series of depolarising and repolarising ion currents mediated by ion channels (Figure 1) [20–22]. Alterations in any of these currents distort the timing and shape of the action potential and leaves the myocardium vulnerable to dysrhythmia. The excitability of the cardiac cell also critically depends on availability of intracellular free calcium; if this is too high, the cell can spontaneously depolarise early (such as seen with digoxin toxicity and CPVT).

### Dysfunctional Ion Channels

The ion channelopathies result from mutations in genes encoding channels or related proteins, altering their properties. A mutation may make a channel non-functional, underactive, overactive or leaky. An example is the cardiac sodium channel Nav1.5 and its encoding gene *SCN5A*; failure of the channel to close results in LQT type 3, and failure to open effectively, or to express functionally, causes Brugada syndrome. The same mutation may cause either within the same family [23].

## Types of Arrhythmia

The cardiac ion channelopathies cause sudden death by causing rapid, usually polymorphic ventricular tachycardia (VT) or ventricular fibrillation (VF). At a tissue level the mechanisms can be *triggered* (due to an after depolarisation such as in LQTS and CPVT) with the arrhythmia sustaining through a circus movement-type re-entry [21], or *re-entry per se* typically due to adjacent areas of the myocardium having different electrophysiological characteristics, such as the right ventricular outflow tract in BrS [24].

## Long QT Syndrome

The term “long QT syndrome” (LQTS) implies the inherited, or genetic, ion channelopathy, whereas “acquired long QT syndrome” is reserved for those where the QT interval is prolonged from acquired heart disease, biochemical or pathophysiological events, such as hypokalaemia, QT-prolonging drugs, hypothermia or neurological events such as a stroke.

The cardinal feature is prolonged repolarisation [14].

### Diagnosis

The diagnosis of LQTS is made clinically, by combining clinical and family history and the 12-lead ECG [25,26], A clinical score (“Schwartz score”) [26] can be a useful guide. Manual measurement of the heart-rate corrected QT interval (QTc), focusses on leads 2 and V5, on two or more ECGs with no extraneous causes of QT prolongation. The “tangent” technique is preferred to determine the end of the T-wave (the steep slope of the T-wave is extended to the baseline) [27]. Use of the Bazett formula (QT length divided by the square root of the preceding R-R interval) is acceptable at all ages [28]. Qualitative assessment of T-wave morphology is as important as QT length (see Figures 1 and 2). Cardiomyopathies often have QT prolongation as part of the disease but the ST segments tend to be depressed inferolaterally and the T-waves biphasic or inverted.

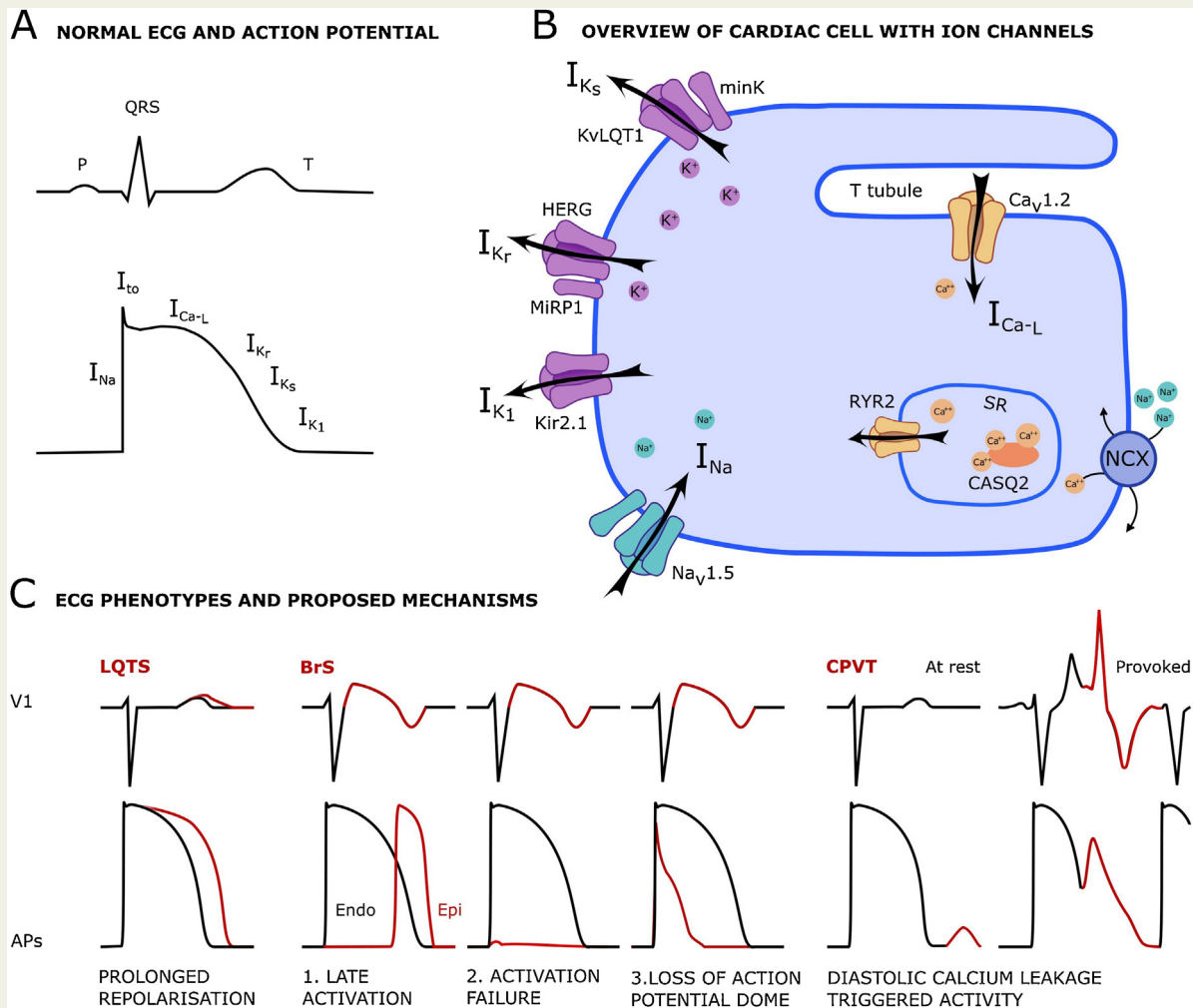
An isolated QTc (in the absence of syncope or family history) >500 ms on more than one occasion is sufficient for the diagnosis of LQTS [25]. Many LQTS subjects have a QTc shorter than this, so a thorough clinical and family history must be taken—something which not all cardiologists do [29]. Following arrhythmic syncope, a diagnosis of LQTS can only be made if a QTc greater than 470 ms on repeated ECGs is found and vasovagal syncope is excluded.

A family history of sudden death can be pivotal to diagnosis [30].

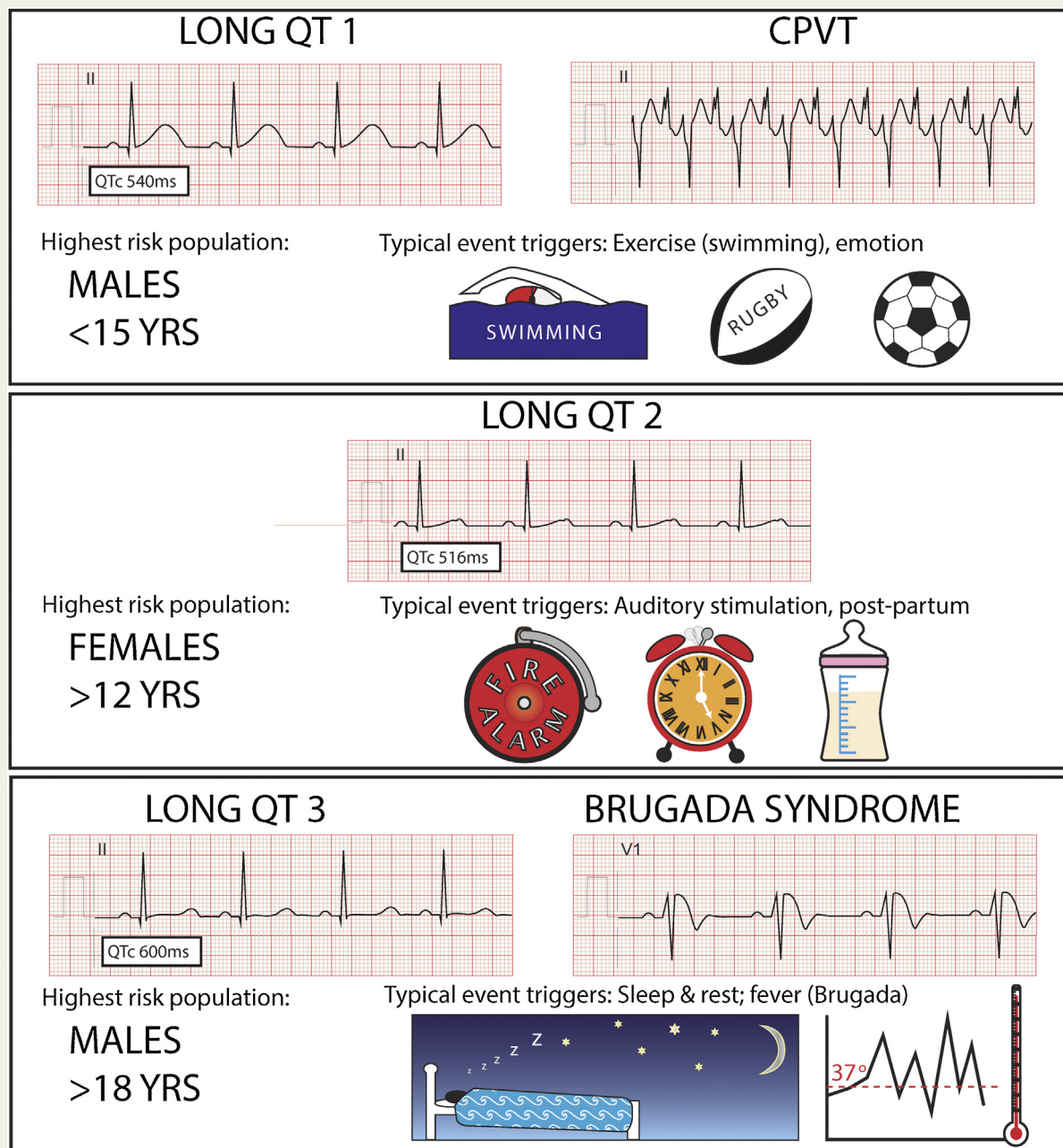
### Ancillary Diagnostic Tests

A QTc >445 ms at 4 minutes in recovery after exercise increases the probability of LQTS, with increased specificity using a cut-off of >480 ms [31,32]. Normative values are available for children [33].

Within 10 seconds of brisk-standing, heart-rate increases [34]. In those affected by LQTS, the QT interval reduces much less than the RR interval. Ventricular ectopic beats or T-wave alternans and T-wave dysmorphology may be seen. This test can be problematic in “fidgety” young children; and



**Figure 1** (A) The cellular action potential driving the heart cycle is shaped by a series of depolarising and repolarising ion currents ( $I$ ). The major depolarising currents in ventricular cardiomyocytes are sodium ( $I_{Na}$ ) and L-type calcium currents ( $I_{Ca-L}$ ). The major repolarising currents are potassium currents (transient outward potassium current  $I_{to}$ , rapid delayed rectifier current  $I_{K_r}$ , and slow delayed rectifier current  $I_{K_s}$ ).  $I_{K_1}$  is an inward rectifier current maintaining resting membrane potential and controlling cellular excitability. (B) The ion channels and related proteins responsible for depolarising ( $Na_v1.5$  and  $Ca_v1.2$ ) and repolarising ( $K_vLQT1/minK$ ,  $HERG/MiRP1$  and  $Kir2.1$ ) currents are found in the cell membrane, either on the cell surface or in the transverse tubules (T tubules). The sodium/calcium exchanger (NCX1) contributes to the depolarising current via changing 3 sodium ions ( $Na^+$ ) for 1 calcium ion ( $Ca^{2+}$ ), resulting in a net positive inward current. Calcium handling and control during cardiomyocyte contraction and relaxation is mediated by the process of calcium-induced calcium release from the sarcoplasmic reticulum (SR), where calcium is bound by calsequestrin (CASQ2) and released into the cytosol by the ryanodine-receptor (RyR2) channel. (C) Suggested mechanisms of the LQTS, Brugada and CPVT ECG patterns, as seen in the first precordial lead (V1). APs: ventricular action potentials. LQTS: Reduced repolarising currents ( $I_{K_s}$  in LQT1 and  $I_{K_r}$  in LQT2) or increased depolarising currents ( $I_{Na}$  in LQT3) result in a prolonged repolarisation and a prolonged QT interval on the ECG. BrS: Three alternative pathophysiological mechanisms underlying the type 1 Brugada pattern have been proposed: (1) late activation of the right ventricle causes ST-segment elevation and repolarisation of the same myocardium causes the negative T-wave, (2) excitation failure at the right ventricular subepicardium causes ST-segment elevation and moderate activation delay at neighbouring sites causes the negative T-wave. (3), loss of the action potential dome at the right ventricular subepicardium but not the subendocardium, i.e. transmural dispersion in action potential duration. CPVT: Resting ECG features in CPVT are typically normal. Dysfunction of the sarcoplasmic reticulum calcium release channel or calcium storage causes leakage of calcium in diastole, and increasing intracellular calcium concentrations causes delayed after-depolarisations and extrasystolic action potentials, that may trigger polymorphic VT. Abbreviations: CPVT, catecholaminergic polymorphic ventricular tachycardia; VT, ventricular tachycardia; ECG, electrocardiogram; LQTS, long QT syndrome.



**Figure 2** Diagram displaying typical phenotypic characteristics of the common cardiac ion channelopathies.

normative values are different from adults; at maximal tachycardia, mean QTc prolongation was 79 ms in children, vs 50 ms in adults [35].

### Genetics

Long QT syndrome is caused by mutations in any of 17 LQTS genes (Table 1) [14,36]. Inheritance is usually autosomal dominant (previously called Romano-Ward syndrome).

The most common genotypes are LQT1, 2, 3 and 5 [37]. In each, a dysfunctional cardiac ion-channel results in prolongation and/or distortion of the cardiac action potential, and thus the QT interval and T-wave. Many of the hundreds

of mutations are unique to a family or very rare. About a quarter of families with LQTS do not yet have a recognised genetic locus.

Two recently described syndromes are very severe and seem to affect only infants and young children: calmodulin related disease (genes, *CALM 1,2 and 3*) associated with seizures and developmental delay [12]; and the so-called triadin-knock out syndrome with recessive inheritance and inverted T-waves [38,39].

Determination of genotype can be useful to confirm diagnosis, allow genetic screening of potentially affected family members, assess degree of risk and hence tailor therapy.



**Table 1** Long QT genes.

Clinical name	Chromosomal locus	Gene name	Current Affected	Non cardiac effects
LQT1	11p15.5	KCNQ1 ( <i>KVLQT1</i> )	K <sup>+</sup> (I <sub>Ks</sub> )	Deafness with recessive form (JLNS)
LQT2	7q35-36	HERG ( <i>KCNH2</i> )	K <sup>+</sup> (I <sub>Kr</sub> )	
LQT3	3p21-24	SCN5A	Na <sup>+</sup> (I <sub>NA</sub> )	
LQT4	4q25-27	Ankyrin B	Na <sup>+</sup> (I <sub>NA</sub> )	
LQT5	21q22.1-22.2	KCNE1 (minK)	K <sup>+</sup> (I <sub>Ks</sub> )	Deafness with recessive form (JLNS)
LQT6	21q22.1-22.2	KCNE2 (MiRP1)	K <sup>+</sup> (I <sub>Kr</sub> )	
LQT7 (Anderson)	17q23	KCNJ2	K <sup>+</sup> (K <sub>ir2.1</sub> )	Anderson-Tahwil syndrome with some mutations
LQT8 (Timothy)	12p13.3	CACNA1C	Ca <sup>2+</sup> + (I <sub>Ca-L</sub> )	Timothy syndrome with some mutations
LQT9	3p25	CAV3 ( <i>Caveolin</i> )	Na <sup>+</sup> (I <sub>NA</sub> )	
LQT10	11q23.3	SCN4B	Na <sup>+</sup> (I <sub>NA</sub> )	
LQT11	7q21-q22	AKAP9 ( <i>A –anchor protein 9</i> )	K <sup>+</sup> (I <sub>Ks</sub> )	
LQT12	20q11.2	SNTA1 ( <i>alpha-1 syntrophin</i> )	Na <sup>+</sup> (I <sub>NA</sub> )	
LQT13	11q24.3	KCNJ5	K <sup>+</sup> (K <sub>ir</sub> )	
LQT14	14q24-q31	Calmodulin1	Many <sup>#</sup>	Seizures, developmental delay
LQT15	2p21.1-p21.3	Calmodulin2	Many <sup>#</sup>	Seizures, developmental delay
LQT16	19q13.2-q13.3	Calmodulin3	Many <sup>#</sup>	Seizures, developmental delay
LQT17		Triadin		

LQT7-Anderson syndrome is a rare neurological disorder characterized by periodic paralysis, skeletal developmental abnormalities, and QT prolongation.

<sup>#</sup>Calmodulin.

### Genotype-Phenotype Correlation in LQTS and Indicators of High Risk

The three most common genotypes (LQT1, LQT2 and LQT3) tend to have genotype-specific syncope triggers for cardiac events, have characteristic T-wave morphologies [40], and age and gender correlated features of high risk (see Figure 2). Subjects with LQT1 and LQT2 tend to have several “warning” syncopal episodes before a sudden death, whereas in LQT3 the first presentation is commonly sudden death [41,42].

In LQT1, boys aged 5–15 years are at highest risk, especially during exercise and particularly swimming, and they have a broad T-wave. With LQT2, it is adult women, particularly up to 9 months post-partum who are at highest risk. Auditory or emotional stimulations feature, and nocturnal events are common, arrhythmias are usually pause-dependent, and the T-wave has a low amplitude notched, or “double-bump”, appearance. With LQT3, gene carriers are often bradycardic with late onset T waves, and sudden death during sleep.

The strongest predictors for high risk are previous cardiac arrest or syncope and a QTc interval recorded at any time during follow-up of over 500 ms [43].

### Multiple Mutations

Approximately 5% of families have two mutations, and family members with both mutations tend to be more severely affected [44]. Two mutations on opposite chromosomes in either the LQT1 or LQT5 gene causes a severe autosomal recessive form of LQTS usually with associated sensorineural deafness, low gastric acid secretion and iron deficiency anaemia (Jervell and Lange-Nielsen syndrome, JLNS) [45].

### Not All Missense Mutations Are the Same

As increasing data are collated, some individual mutations seem to be rather benign and others more malignant [46]. In general, adult women with LQT2 are at greater risk of cardiac arrest than men. However if the missense mutation is in the pore loop regions, men are at greater risk [47].

### Gene Modifiers

Modifiers of gene expression include untranslated regions which modify the RNA binding site in LQT1 [32] and minor changes (single nucleotide polymorphisms) in NOS1AP, a gene linked to QT prolongation in the general population [33].

LQT8-Timothy syndrome is a rare condition characterised by syndactyly, facial dysmorphism, autism and severe LQTS.

The polymorphisms S1103Y in SCN5A in Blacks, and D85N in KCNE1 in Whites, increase the chance of medication or drug-induced QT prolongation and arrhythmia [48]. They do so by causing minor dysfunction, giving a reduced repolarisation reserve which is only unmasked by a drug which exacerbates the dysfunction [15].

### Therapy

There are four levels of therapy in LQTS, applied according to severity.

1. All gene carriers must avoid QT prolonging drugs (see [www.crediblemeds.org](http://www.crediblemeds.org)). Caution when swimming, especially for LQT1; remove loud alarm clocks for LQT2.

2. Long acting beta blockers are recommended in all types, including LQT3 [46]; and, are given to most except those at the lowest risk, such as asymptomatic pre-pubertal females and adults over 20 years with LQT1 and a consistently normal QT interval [49].

3. Left cardiac sympathetic denervation is especially effective in (high risk) LQT1, or in those who need but are unable to take beta blockers. Side-effects are common but usually well tolerated [50].

4. Implanted defibrillators are reserved for those who have had a cardiac arrest, those with syncope whilst taking beta blockers and others still at high risk despite the above therapies [51,52], such as adult women with LQT2 and a QTc over 550 ms especially if there is a history of syncope [14,53].

## J-Wave Syndromes: Brugada Syndrome and Early Repolarisation Syndrome

The J-wave syndromes include BrS and early repolarisation syndrome (ERPS) [60]. They are distinct entities but have some clinical similarities, namely they share a similar predisposition for sudden cardiac death in the third decade of life, male predominance and response to medical therapy [60]. The pathophysiological mechanism of these disorders is disputed (see Figure 1) [60]. The mechanism of arrhythmogenesis may be quite distinct as well [54,55].

### Brugada Syndrome

Brugada syndrome is diagnosed when the typical ECG signature is observed in an apparent structurally normal heart and is more prevalent in South Asian countries. This ECG signature associates with a risk of potentially lethal arrhythmias that typically occur under vagal triggers (e.g., after meals and during the night) or during fever [18]. Risk of arrhythmias is highest in the fourth and fifth decades, and males are more affected than females.

The pathophysiology of the ECG features and the arrhythmogenic substrate are disputed but increasing evidence is emerging that minor structural abnormalities in the right ventricular outflow tract (RVOT) area underlie the disease. As such, BrS may have to be regarded as part of the ARVC spectrum (arrhythmogenic right ventricular cardiomyopathy) [56].

#### Diagnosis

To diagnose BrS, the typical 'type 1' ECG is required (Figures 1 and 2) [57], where right precordial concave ST segment elevation (J-point elevation of at least 2 mm), followed by ST-depression, occurs in the same leads. The ECG sign may not be consistently present and could be unmasked by a class Ic drug challenge (i.e. ajmaline, flecainide). *However, when the ECG is unmasked by drugs, additional clinical criteria are required for the diagnosis BrS* [57]. Associated arrhythmias have their origin in the RVOT; usually, a short coupled extra systole is followed by a rapid polymorphic VT/VF.

### Genetics

The genetics of BrS is more complex than the other primary arrhythmia syndromes. The first gene identified, i.e. *SCN5A*, still stands as a potential causal gene but all other 20+ genes, encoding for different ion channel (subunits) are now all disputed as causal, monogenetic, causes for BrS [58]. In fact, there is increasing evidence that BrS is an oligogenetic disease, with

involvement of more than one genetic factor with different effect sizes [59]. The more of these genetic factors one has, the higher the likelihood of having a type 1 Brugada pattern.

Currently, molecular genetic testing should be limited to *SCN5A* and in *SCN5A* families (presymptomatic) and counselling should include an ECG, because phenotype positive-genotype negative cases have been described within these families [60]. There might be some role for genetic testing in risk stratification [61].

#### Therapy

All patients are advised to manage fevers and avoid large meals late before going to bed. An implantable cardiac defibrillator (ICD) is indicated for resuscitated patients, and those with documented ventricular arrhythmias or typical arrhythmic syncope. The risk for asymptomatic patients with a spontaneous type 1 ECG is not well defined and methods to define their risk more accurately are disputed. Different studies provide controversial results [62]. Probably the most reliable risk markers are spontaneous variation in the ECG pattern and marked fractionation of the QRS complex.

An emerging therapy is ablation of fractionated signals picked up from the epicardial layer of the RVOT. Extinction of all fractionated activity in this area normalises the ECG and abolishes arrhythmias during initially reported follow-up [63]. Pharmacological therapies exist in the form of oral quinidine or, in acute situations with an arrhythmic storm, isoprenaline given intravenously.

### ERPS (Early Repolarisation Syndrome)

Elevation of the J point, at the end of the QRS complex is a common, benign phenomenon, occurring in the inferolateral ECG leads. It is diagnosed when a J point notch or ST segment is elevated greater than 1 mm in two or more contiguous inferior and or anterior leads excluding V<sub>1</sub>-V<sub>3</sub> [25].

However, early repolarisation is more prevalent among those with idiopathic VF, and rare rhythm strips have shown a rising J point prior to VF onset [64]. However, the ECG appearance of ERPS is so common (over 5% of the normal population, and 25% of athletes), that it cannot be used as a screening tool for risk of sudden death on a population basis. When identified after a cardiac arrest or an autopsy negative sudden death, and other causes are excluded, it may be diagnosed as a cause of the event if ECG criteria are met [57].

Like in BrS, arrhythmia prevention may be achieved with quinidine along with defibrillator back-up, and arrhythmia storms can be managed with isoprenaline [57].

Genetic testing has no value in these patients to date.

### CPVT (Catecholaminergic Polymorphic Ventricular Tachycardia)

#### Typical Presentation and Diagnosis

The typical presentation is a child between the age of 4 and 12 years presenting with sudden exercise-related syncope or cardiac arrest, often related to swimming, and tending to be worse in males [65,66]. Sudden Arrhythmia Death

Syndrome autopsy series find CPVT almost as commonly as long QT syndrome [7,67], so, given it is much rarer than LQTS, it is clearly much more severe. Cases in infancy (sudden infant death) are rare, and milder or later presenting forms in mid-adult life are being recognised increasingly. Many syncopal episodes in fact occur during “wakeful rest” [66]. The resting 12-lead ECG is normal. The diagnosis is made by exercise testing, after the exclusion of structural heart disease, by documenting premature ventricular contractions usually at heart rates over 100 beats per minute on exercise testing, which progress to polymorphic VT, and sometimes to the classic “bidirectional VT” which is pathognomonic (see Figure 1).

### Cellular Basis and Genetics

Catecholaminergic polymorphic VT mutations lead to increased calcium release from the sarcoplasmic reticulum during diastole, and adrenaline stimulates further calcium release, resulting in delayed after depolarisations and triggered activity (see Figure 1). Approximately 60% of patients with CPVT have a mutation in the cardiac ryanodine receptor gene *RyR2* (CPVT1) [66,68]. Although this is autosomal dominant, cases presenting in young childhood are commonly *de novo*, reflecting the severity of this condition. The mutations are highly penetrant. CPVT2 is autosomal recessive and very uncommon, caused by calsequestrin mutations (*CASQ2*) [69]. Other rare cases have been linked to other calcium handling genes *CALM1* (encoding calmodulin) [70] and *TRDN* (encoding Triadin) [38]. *KCNJ2* mutations and *TECRL* have also been implicated [71,72].

### Therapies and Risk Stratification

The mainstay of therapy is beta blockade. However breakthrough events are common [65]. There have been three significant lessons relating to management over the last decade:

1. Left cardiac sympathectomy was shown to be effective [73].
2. Flecainide prevents VT by inhibiting RyR2-mediated calcium release [74,75]. If beta blockers cannot be tolerated, flecainide has been used alone [76]. It also works in RyR2-negative cases [77].
3. It was recognised that ICDs can be counter-productive in polymorphic VT due to CPVT; a shock may be ineffective, and the ensuing adrenergic output can worsen the VT storm [78]. Guidelines currently suggest ICDs should be implanted after cardiac arrest for CPVT [79]; although, this is being questioned due to the remarkable efficacy of flecainide, beta blockade and cardiac sympathectomy [80]. Implantable cardiac defibrillators should be programmed to shock after a long delay, to cardiovert VF rather than worsen the VT [78].

### On the Horizon for Cardiac Ion Channelopathies

Being largely monogenetic conditions, there is the possibility of gene therapy. A mouse *CASQ2* model has been created and then cured by viral transfection [81]. We might hope that pluripotent cell lines might lead to new therapies in

cardiac ion channelopathies, such as flecainide has become for CPVT [82]. In the meantime, clinicians have a responsibility to screen families of individuals with a channelopathy, or an unexplained sudden death, and encourage families to participate in clinical registries to further the goal of individualised therapy.

## References

- [1] Bagnall RD, Weintraub RG, Ingles J, Duflou J, Yeates L, Lam L, et al. A prospective study of sudden cardiac death among children and young adults. *N Engl J Med* 2016;374(25):2441–52.
- [2] Behr ER, Dalageorgou C, Christiansen M, Syrris P, Hughes S, Tome Esteban MT, et al. Sudden arrhythmic death syndrome: familial evaluation identifies inheritable heart disease in the majority of families. *Eur Heart J* 2008;29(13):1670–80.
- [3] Tan HL, Hofman N, van Langen IM, van der Wal AC, Wilde AA. Sudden unexplained death: heritability and diagnostic yield of cardiological and genetic examination in surviving relatives. *Circulation* 2005;(July):207–13. 2005.
- [4] Skinner JR, Crawford J, Smith W, Aitken A, Heaven D, Evans CA, et al. Prospective, population-based long QT molecular autopsy study of post-mortem negative sudden death in 1 to 40 year olds. *Heart Rhythm* 2011;8(3):412–9.
- [5] Skinner JR, Duflou JA, Semsarian C. Reducing sudden death in young people in Australia and New Zealand: the TRAGADY initiative. *Med J Aust* 2008;189(10):539–40.
- [6] Tester DJ, Medeiros-Domingo A, Will ML, Haglund CM, Ackerman MJ. Cardiac channel molecular autopsy: insights from 173 consecutive cases of autopsy-negative sudden unexplained death referred for postmortem genetic testing. *Mayo Clin Proc* 2012;87(6):524–39.
- [7] Lahrouchi N, Raju H, Lodder EM, Papatheodorou E, Ware JS, Papadakis M, et al. Utility of post-mortem genetic testing in cases of sudden arrhythmic death syndrome. *J Am Coll Cardiol* 2017;69(17):2134–45.
- [8] Risgaard B, Winkel BG, Jabbari R, Glinge C, Ingemann-Hansen O, Thomsen JL, et al. Sports-related sudden cardiac death in a competitive and a noncompetitive athlete population aged 12 to 49 years: data from an unselected nationwide study in Denmark. *Heart Rhythm* 2014;11(10):1673–81.
- [9] Finocchiaro G, Papadakis M, Robertus JL, Dhutia H, Steriotis AK, Tome M, et al. Etiology of sudden death in sports: insights from a United Kingdom Regional Registry. *J Am Coll Cardiol* 2016;67(18):2108–15.
- [10] Maron BJ, Haas TS, Ahluwalia A, Murphy CJ, Garberich RF. Demographics epidemiology of sudden deaths in young competitive athletes: from the United States National Registry. *Am J Med* 2016;129(11):1170–7.
- [11] Goldman AM, Behr ER, Semsarian C, Bagnall RD, Sisodiya S, Cooper PN. Sudden unexpected death in epilepsy genetics: molecular diagnostics and prevention. *Epilepsia* 2016;57(Suppl. 1):17–25.
- [12] Crotti L, Johnson CN, Graf E, De Ferrari GM, Cuneo BF, Ovadia M, et al. Calmodulin mutations associated with recurrent cardiac arrest in infants. *Circulation* 2013;127(9):1009–17.
- [13] MacCormick JM, McAlister H, Crawford J, French JK, Crozier I, Shelling AN, et al. Misdiagnosis of long QT syndrome as epilepsy at first presentation. *Ann Emerg Med* 2009;54(1):26–32.
- [14] Waddell-Smith KE, Skinner JR, members of the CSANZ Genetics Council Writing Group. Update on the diagnosis and management of familial long QT syndrome. *Heart Lung Circ* 2016;25(8):769–76.
- [15] Roden DM. Predicting drug-induced QT prolongation and torsades de pointes. *J Physiol* 2016;594(9):2459–68.
- [16] Probst V, Denjoy I, Meregalli PG, Amiraault JC, Sacher F, Mansourati J, et al. Clinical aspects and prognosis of Brugada syndrome in children. *Circulation* 2007;115(15):2042–8.
- [17] Schwartz PJ, Stramba-Badiale M, Crotti L, Pedrazzini M, Besana A, Bosi G, et al. Prevalence of the congenital long-QT syndrome. *Circulation* 2009;120(18):1761–7.
- [18] Mizusawa Y, Wilde AA. Brugada syndrome. *Circ Arrhythm Electrophysiol* 2012;5(3):606–16.
- [19] Probst V, Veltmann C, Eckardt L, Meregalli PG, Gaita F, Tan HL, et al. Long-term prognosis of patients diagnosed with Brugada syndrome: results from the FINGER Brugada Syndrome Registry. *Circulation* 2010;121(5):635–43.



- [20] Abriel H. Cardiac sodium channel Na(v)1.5 and interacting proteins: Physiology and pathophysiology. *J Mol Cell Cardiol* 2010;48(1):2–11.
- [21] Tse G, Chan YW, Keung W, Yan BP. Electrophysiological mechanisms of long and short QT syndromes. *Int J Cardiol Heart Vasc* 2017;14:8–13.
- [22] Bohnen MS, Peng G, Robey SH, Terrenoire C, Iyer V, Sampson KJ, et al. Molecular pathophysiology of congenital long QT syndrome. *Physiol Rev* 2017;97(1):89–134.
- [23] Remme CA, Wilde AA, Bezzina CR. Cardiac sodium channel overlap syndromes: different faces of SCN5A mutations. *Trends Cardiovasc Med* 2008;18(3):78–87.
- [24] Martin CA, Matthews GD, Huang CL. Sudden cardiac death and inherited channelopathy: the basic electrophysiology of the myocyte and myocardium in ion channel disease. *Heart* 2012;98(7):536–43.
- [25] Priori SG, Wilde AA, Horie M, Cho Y, Behr ER, Berul C, et al. HRS/EHRA/APHRS Expert Consensus Statement on the Diagnosis and Management of Patients with Inherited Primary Arrhythmia Syndromes: Document endorsed by HRS, EHRA, and APHRS in May 2013 and by ACCF, AHA, PACES, and AEPCC in June 2013. *Heart Rhythm* 2013;10(12):1932–63.
- [26] Schwartz PJ, Moss AJ, Vincent GM, Crampton RS. Diagnostic criteria for the long QT syndrome. An update. *Circulation* 1993;88(2):782–4.
- [27] Waddell-Smith K, Gow RM, Skinner JR. How to measure a QT interval. *Med J Aust* 2017;207(3):107–10.
- [28] Phan DQ, Silka MJ, Lan YT, Chang RK. Comparison of formulas for calculation of the corrected QT interval in infants and young children. *J Pediatr* 2015;166(4): 960–4.e1–2.
- [29] Waddell-Smith KE, Donoghue T, Oates S, Graham A, Crawford J, Stiles MK, et al. Inpatient detection of cardiac-inherited disease: the impact of improving family history taking. *Open Heart* 2016;3(1):e000329.
- [30] Colman N, Bakker A, Linzer M, Reitsma JB, Wieling W, Wilde AA. Value of history-taking in syncope patients: in whom to suspect long QT syndrome? *Europace* 2009;11(7):937–43.
- [31] Chattha IS, Sy RW, Yee R, Gula LJ, Skanes AC, Klein GJ, et al. Utility of the recovery electrocardiogram after exercise: a novel indicator for the diagnosis and genotyping of long QT syndrome? *Heart Rhythm* 2010;7(7):906–11.
- [32] Sy RW, van der Werf C, Chattha IS, Chockalingam P, Adler A, Healey JS, et al. Derivation and validation of a simple exercise-based algorithm for prediction of genetic testing in relatives of LQTS probands. *Circulation* 2011;124(20):2187–94.
- [33] Berger WR, Gow RM, Kambari S, Cheung M, Smith KR, Davis AM. The QT and corrected QT interval in recovery after exercise in children. *Circ Arrhythm Electrophysiol* 2011;4(4):448–55.
- [34] Viskin S, Postema PG, Bhuiyan ZA, Rosso R, Kalman JM, Vohra JK, et al. The response of the QT interval to the brief tachycardia provoked by standing: a bedside test for diagnosing long QT syndrome. *J Am Coll Cardiol* 2010;55(18):1955–61.
- [35] Filippini L, Postema PG, Zoubin K, Hermans BJM, Blom NA, Delhaas T, et al. The brisk-standing-test for long QT syndrome in prepubertal school children: defining normal. *Europace* 2018;20(F1):f108–12.
- [36] Modell SM, Lehmann MH. The long QT syndrome family of cardiac ion channelopathies: a HuGE review. *Genet Med* 2006;8(3):143–55.
- [37] Earle N, Crawford J, Smith W, Hayes I, Shelling A, Hood M, et al. Community detection of long QT syndrome with a clinical registry: an alternative to ECG screening programs? *Heart Rhythm* 2013;10(2):233–8.
- [38] Roux-Buisson N, Cacheux M, Fourest-Lieuvain A, Fauconnier J, Brocard J, Denjoy I, et al. Absence of triadin, a protein of the calcium release complex, is responsible for cardiac arrhythmia with sudden death in human. *Hum Mol Genet* 2012;21(12):2759–67.
- [39] Altmann HM, Tester DJ, Will ML, Middha S, Evans JM, Eckloff BW, et al. Homozygous/compound heterozygous triadin mutations associated with autosomal-recessive long-QT syndrome and pediatric sudden cardiac arrest: elucidation of the triadin knockout syndrome. *Circulation* 2015;131(23):2051–60.
- [40] Schwartz PJ, Priori SG, Spazzolini C, Moss AJ, Vincent GM, Napolitano C, et al. Genotype-phenotype correlation in the long-QT syndrome: gene-specific triggers for life-threatening arrhythmias. *Circulation* 2001;103(1):89–95.
- [41] Priori SG, Schwartz PJ, Napolitano C, Bloise R, Ronchetti E, Grillo M, et al. Risk stratification in the long-QT syndrome. *N Engl J Med* 2003;348(19):1866–74.
- [42] Moss AJ, Goldenberg I. Importance of knowing the genotype and the specific mutation when managing patients with long QT syndrome. *Circ Arrhythm Electrophysiol* 2008;1:213–26.
- [43] Sauer AJ, Moss AJ, McNitt S, Peterson DR, Zareba W, Robinson JL, et al. Long QT syndrome in adults. *J Am Coll Cardiol* 2007;49(3):329–37.
- [44] Mullally J, Goldenberg I, Moss AJ, Lopes CM, Ackerman MJ, Zareba W, et al. Risk of life-threatening cardiac events among patients with long QT syndrome and multiple mutations. *Heart Rhythm* 2012;10(3):378–82.
- [45] Winbo A, Sandstrom O, Palmqvist R, Rydberg A. Iron-deficiency anaemia, gastric hyperplasia, and elevated gastrin levels due to potassium channel dysfunction in the Jervell and Lange-Nielsen Syndrome. *Cardiol Young* 2012;1–10.
- [46] Wilde AA, Moss AJ, Kaufman ES, Shimizu W, Peterson DR, Benhorin J, et al. Clinical aspects of type 3 Long-QT syndrome: an International Multicenter Study. *Circulation* 2016;134(12):872–82.
- [47] Migdalovich D, Moss AJ, Lopes CM, Costa J, Ouellet G, Barsheshet A, et al. Mutation and gender-specific risk in type 2 long QT syndrome: implications for risk stratification for life-threatening cardiac events in patients with long QT syndrome. *Heart Rhythm* 2011;8(10):1537–43.
- [48] Splawski I, Timothy KW, Tateyama M, Clancy CE, Malhotra A, Beggs AH, et al. Variant of SCN5A sodium channel implicated in risk of cardiac arrhythmia. *Science* 2002;297(5585):1333–6.
- [49] Waddell-Smith KE, Earle N, Skinner JR. Must every child with long QT syndrome take a beta blocker? *Arch Dis Child* 2015;100(3):279–82.
- [50] Waddell-Smith KE, Ertresvaag KN, Li J, Chaudhuri K, Crawford JR, Hamill JK, et al. Physical and psychological consequences of left cardiac sympathetic denervation in long-QT syndrome and catecholaminergic polymorphic ventricular tachycardia. *Circ Arrhythm Electrophysiol* 2015;8(5):1151–8.
- [51] Gaba P, Bos JM, Cannon BC, Cha YM, Friedman PA, Asirvatham SJ, et al. Implantable cardioverter-defibrillator explantation for overdiagnosed or overtreated congenital long QT syndrome. *Heart Rhythm* 2016;13(4):879–85.
- [52] Schwartz PJ, Spazzolini C, Priori SG, Crotti L, Vicentini A, Landolina M, et al. Who are the long-QT syndrome patients who receive an implantable cardioverter-defibrillator and what happens to them?: data from the European Long-QT Syndrome Implantable Cardioverter-Defibrillator (LQTS ICD) Registry. *Circulation* 2010;122(13):1272–82.
- [53] Horner JM, Kinoshita M, Webster TL, Haglund CM, Friedman PA, Ackerman MJ. Implantable cardioverter defibrillator therapy for congenital long QT syndrome: a single-center experience. *Heart Rhythm* 2010;7(11):1616–22.
- [54] Junttila MJ, Sager SJ, Tikkanen JT, Anttonen O, Huikuri HV, Myerburg RJ. Clinical significance of variants of J-points and J-waves: early repolarization patterns and risk. *Eur Heart J* 2012;33(21):2639–43.
- [55] Morita H, Zipes DP, Wu J. Brugada syndrome: insights of ST elevation, arrhythmogenicity, and risk stratification from experimental observations. *Heart Rhythm* 2009;6(11 Suppl):S34–43.
- [56] Corrado D, Zorzi A, Cerrone M, Rigato I, Mongillo M, Bauce B, et al. Relationship between arrhythmogenic right ventricular cardiomyopathy and Brugada syndrome: new insights from molecular biology and clinical implications. *Circ Arrhythm Electrophysiol* 2016;9(4):e003631.
- [57] Antzelevitch C, Yan GX, Ackerman MJ, Borggreffe M, Corrado D, Guo J, et al. J-Wave syndromes expert consensus conference report: emerging concepts and gaps in knowledge. *Europace* 2017;19(4):665–94.
- [58] Le Scouarnec S, Karakachoff M, Gourraud JB, Lindenbaum P, Bonnaud S, Portero V, et al. Testing the burden of rare variation in arrhythmia-susceptibility genes provides new insights into molecular diagnosis for Brugada syndrome. *Hum Mol Genet* 2015;24(10):2757–63.
- [59] Bezzina CR, Barc J, Mizusawa Y, Remme CA, Gourraud JB, Simonet F, et al. Common variants at SCN5A-SCN10A and HEY2 are associated with Brugada syndrome, a rare disease with high risk of sudden cardiac death. *Nat Genet* 2013;45(9):1044–9.
- [60] Probst V, Wilde AA, Barc J, Sacher F, Babuty D, Mabo P, et al. SCN5A mutations and the role of genetic background in the pathophysiology of Brugada syndrome. *Circ Cardiovasc Genet* 2009;2(6):552–7.
- [61] Yamagata K, Horie M, Aiba T, Ogawa S, Aizawa Y, Ohe T, et al. Genotype-phenotype correlation of SCN5A mutation for the clinical and electrocardiographic characteristics of probands with Brugada syndrome: a Japanese Multicenter Registry. *Circulation* 2017;135(23):2255–70.
- [62] Adler A, Rosso R, Chorin E, Havakuk O, Antzelevitch C, Viskin S. Risk stratification in Brugada syndrome: clinical characteristics, electrocardiographic parameters, and auxiliary testing. *Heart Rhythm* 2016;13(1):299–310.
- [63] Nademanee K, Veerakul G, Chandanamattha P, Chaotawee L, Ariyachaipanich A, Jirasirojanakorn K, et al. Prevention of ventricular fibrillation episodes in Brugada syndrome by catheter ablation over the anterior right ventricular outflow tract epicardium. *Circulation* 2011;123(12):1270–9.



- [64] Haissaguerre M, Derval N, Sacher F, Jesel L, Deisenhofer I, de Roy L, et al. Sudden cardiac arrest associated with early repolarization. *N Engl J Med* 2008;358(19):2016–23.
- [65] Leenhardt A, Lucet V, Denjoy I, Grau F, Ngoc DD, Coumel P. Catecholaminergic polymorphic ventricular tachycardia in children. A 7-year follow-up of 21 patients. *Circulation* 1995;91(5):1512–9.
- [66] Roston TM, Yuchi Z, Kannankeril PJ, Hathaway J, Vinocur JM, Etheridge SP, et al. The clinical and genetic spectrum of catecholaminergic polymorphic ventricular tachycardia: findings from an international multi-centre registry. *Europace* 2018;20(3):541–7.
- [67] Semsarian C, Bagnall RD. Sudden cardiac death in children and young adults. *N Engl J Med* 2016;375(13):1301–2.
- [68] Priori SG, Napolitano C, Tiso N, Memmi M, Vignati G, Bloise R, et al. Mutations in the cardiac ryanodine receptor gene (hRyR2) underlie catecholaminergic polymorphic ventricular tachycardia. *Circulation* 2001;103(2):196–200.
- [69] Lahat H, Pras E, Olender T, Avidan N, Ben-Asher E, Man O, et al. A missense mutation in a highly conserved region of CASQ2 is associated with autosomal recessive catecholamine-induced polymorphic ventricular tachycardia in Bedouin families from Israel. *Am J Hum Genet* 2001;69(6):1378–84.
- [70] Nyegaard M, Overgaard MT, Sondergaard MT, Vranas M, Behr ER, Hildebrandt LL, et al. Mutations in calmodulin cause ventricular tachycardia and sudden cardiac death. *Am J Hum Genet* 2012;91(4):703–12.
- [71] Tester DJ, Arya P, Will M, Haglund CM, Farley AL, Makielski JC, et al. Genotypic heterogeneity and phenotypic mimicry among unrelated patients referred for catecholaminergic polymorphic ventricular tachycardia genetic testing. *Heart Rhythm* 2006;3(7):800–5.
- [72] Devalla HD, Gelinas R, Aburawi EH, Beqqali A, Goyette P, Freund C, et al. TECRL, a new life-threatening inherited arrhythmia gene associated with overlapping clinical features of both LQTS and CPVT. *EMBO Mol Med* 2016;8(12):1390–408.
- [73] Wilde AA, Bhuiyan ZA, Crotti L, Facchini M, De Ferrari GM, Paul T, et al. Left cardiac sympathetic denervation for catecholaminergic polymorphic ventricular tachycardia. *N Engl J Med* 2008;358(19):2024–9.
- [74] van der Werf C, Kannankeril PJ, Sacher F, Krahn AD, Viskin S, Leenhardt A, et al. Flecainide therapy reduces exercise-induced ventricular arrhythmias in patients with catecholaminergic polymorphic ventricular tachycardia. *J Am Coll Cardiol* 2011;57(22):2244–54.
- [75] Watanabe H, Chopra N, Laver D, Hwang HS, Davies SS, Roach DE, et al. Flecainide prevents catecholaminergic polymorphic ventricular tachycardia in mice and humans. *Nat Med* 2009;15(4):380–3.
- [76] Padfield GJ, AlAhmari L, Lieve KV, AlAhmari T, Roston TM, Wilde AA, et al. Flecainide monotherapy is an option for selected patients with catecholaminergic polymorphic ventricular tachycardia intolerant of beta-blockade. *Heart Rhythm* 2016;13(2):609–13.
- [77] Watanabe H, van der Werf C, Roses-Noguer F, Adler A, Sumitomo N, Veltmann C, et al. Effects of flecainide on exercise-induced ventricular arrhythmias and recurrences in genotype-negative patients with catecholaminergic polymorphic ventricular tachycardia. *Heart Rhythm* 2013;10(4):542–7.
- [78] Roses-Noguer F, Jarman JW, Clague JR, Till J. Outcomes of defibrillator therapy in catecholaminergic polymorphic ventricular tachycardia. *Heart Rhythm* 2014;11(1):58–66.
- [79] Pflaumer A, Davis AM. Guidelines for the diagnosis and management of catecholaminergic polymorphic ventricular tachycardia. *Heart Lung Circ* 2012;21(2):96–100.
- [80] Pflaumer A. Catecholaminergic polymorphic tachycardia: underestimated and overtreated? *Heart* 2017;103(12):889–90.
- [81] Denegri M, Avelino-Cruz JE, Boncompagni S, De Simone SA, Auricchio A, Villani L, et al. Viral gene transfer rescues arrhythmogenic phenotype and ultrastructural abnormalities in adult calsequestrin-null mice with inherited arrhythmias. *Circ Res* 2012;110(5):663–8.
- [82] Mehta A, Ramachandra CJA, Singh P, Chitre A, Lua CH, Mura M, et al. Identification of a targeted and testable antiarrhythmic therapy for long-QT syndrome type 2 using a patient-specific cellular model. *Eur Heart J* 2018;39(16):1446–55.