

# CARDIAC CHANNELOPATHIES

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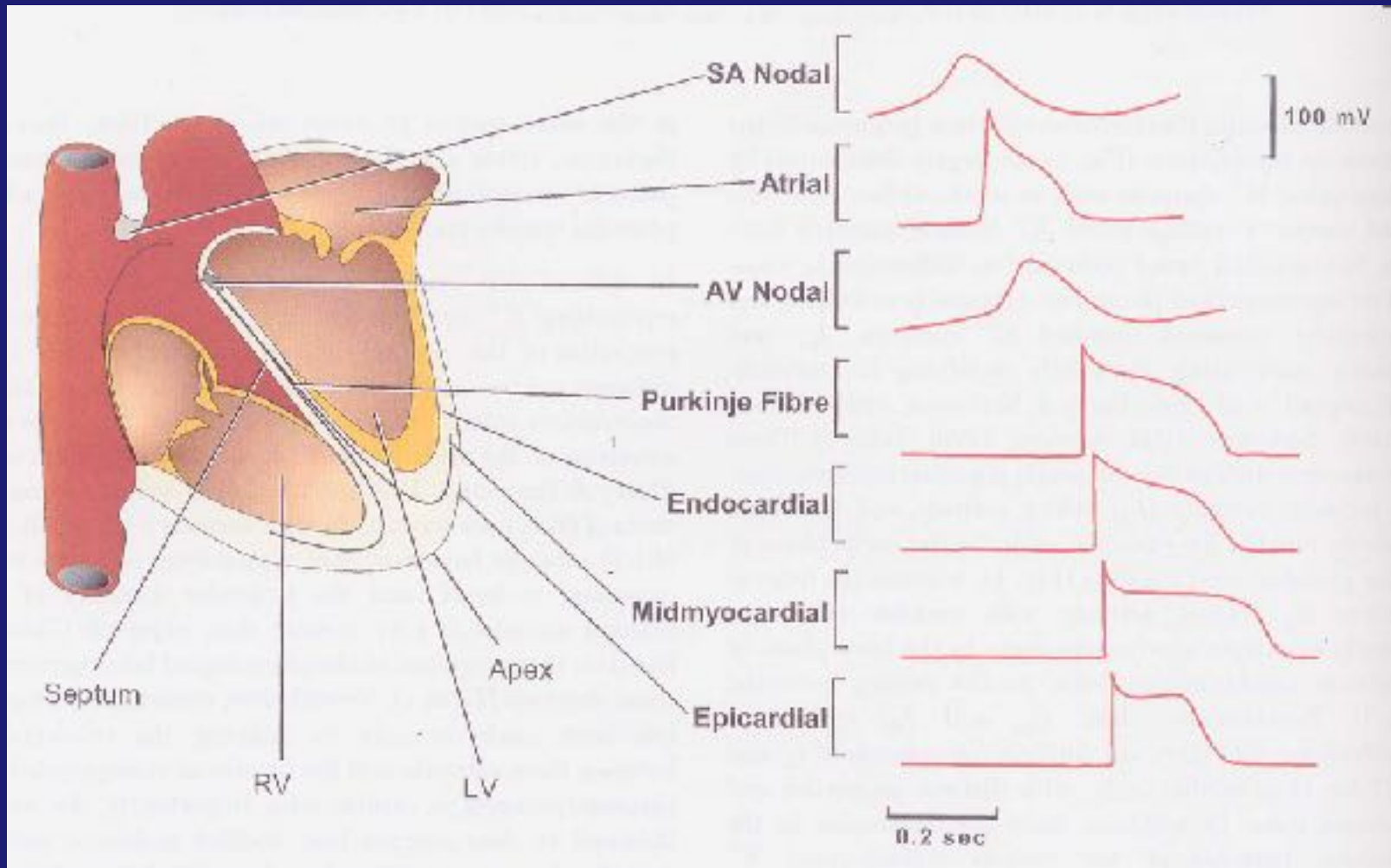
# Outline

- ❖ Introduction
- ❖ Overview of cardiac action potential
- ❖ Structure of ion channels
- ❖ Long QT Syndrome
- ❖ Short QT Syndrome
- ❖ Brugada Syndrome
- ❖ Catecholaminergic Polymorphic Ventricular Tachycardia (CPVT)
- ❖ Idiopathic ventricular fibrillation
- ❖ Progressive cardiac conduction defect
- ❖ Ankyrin-B Syndrome

# Introduction

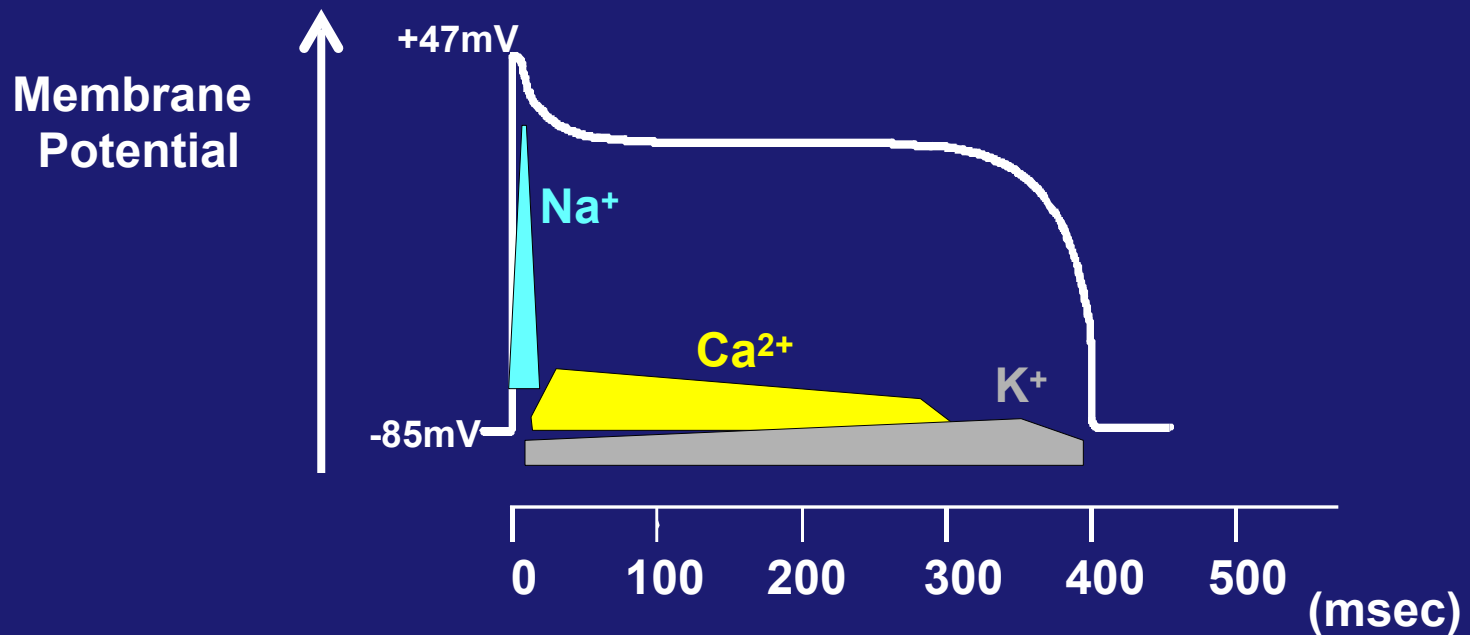
- Cardiac arrhythmias and conduction defects result from abnormalities in three main families of proteins:
  - Contractile proteins (e.g. HOCM)
  - Cytoskeletal proteins (eg. DCM)
  - Ion channels and their regulators

# Action Potential Waveforms in Different Regions of the Heart



Nerbonne JM. J of Physiology 2000, 525.2;285

# The Cardiac Action Potential



# Structure of Ion Channels

**ION  
CHANNEL**

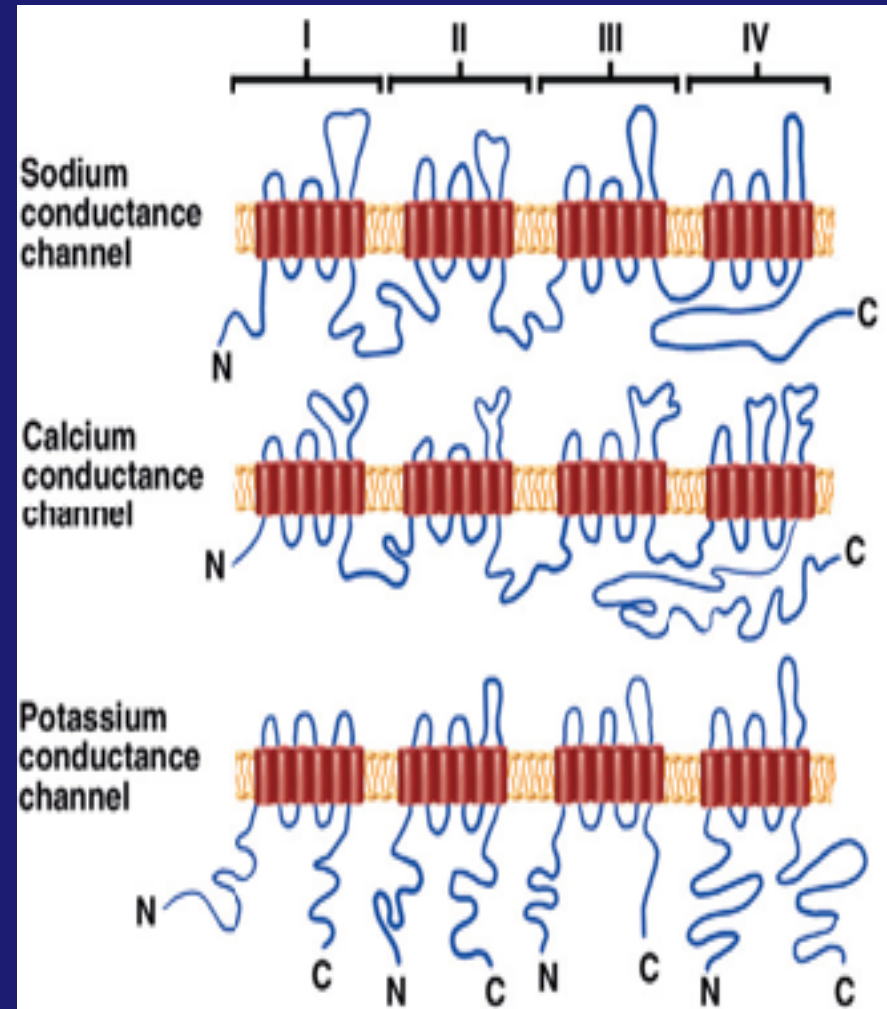


**Alpha Subunit  
(pore forming)**

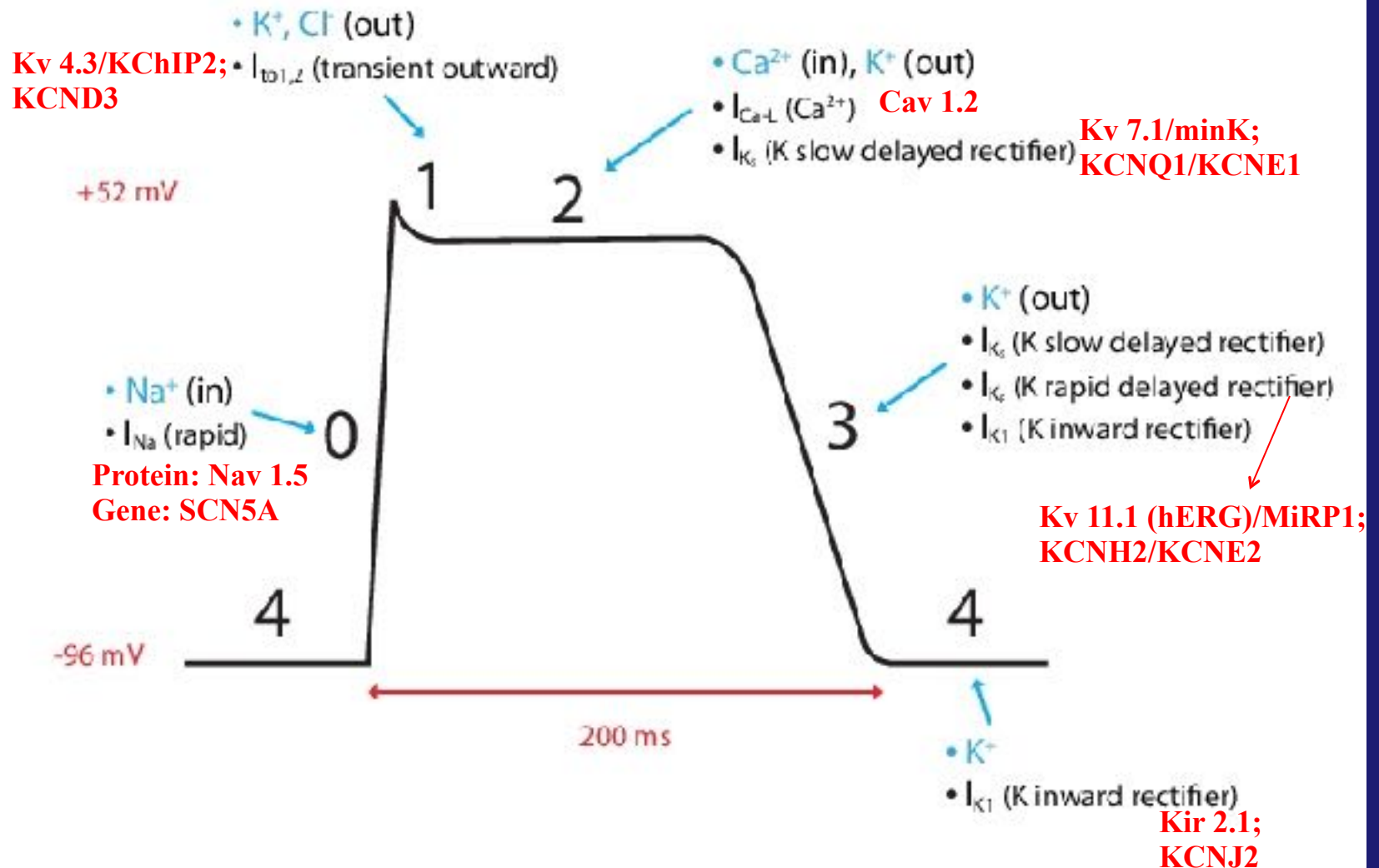


**Auxillary  
subunits ( $\beta, \gamma, \delta$ )**

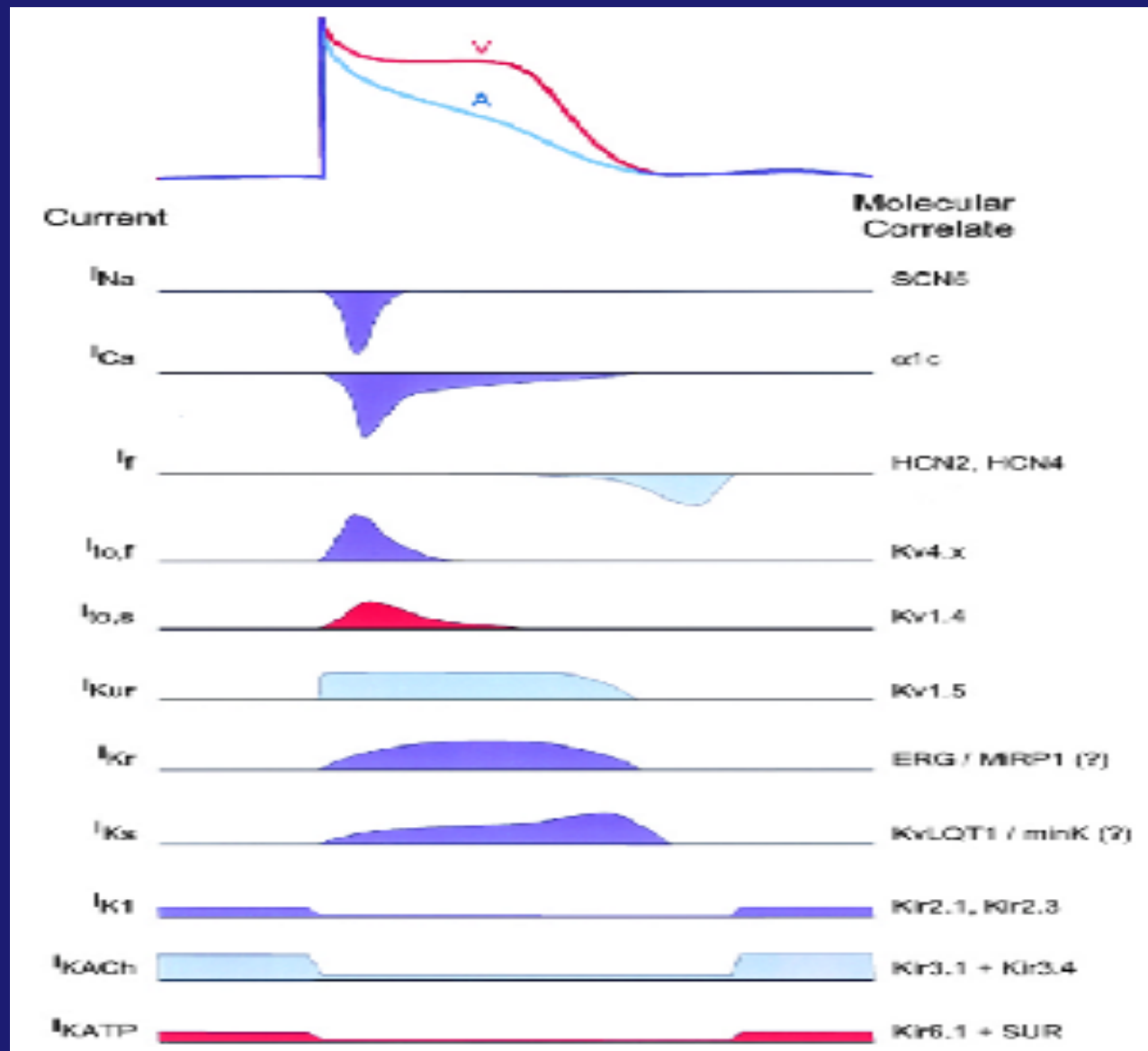
- Voltage-gated  $\text{Na}^+$  and  $\text{Ca}^{2+}$  channels:
  - single tetramer
  - four covalently linked repeats of the six transmembrane-spanning motifs.
- Voltage-gated  $\text{K}^+$  channels:
  - four separate subunits,
  - each containing a single six transmembrane-spanning motif.
- Inwardly rectifying  $\text{K}^+$  channels
  - In contrast to voltage-gated  $\text{K}^+$  channel alpha subunits, the Kir alpha subunits have only two (not six) transmembrane domains.



# Relationship Between Cardiac Action Potential and Ion Channel Currents

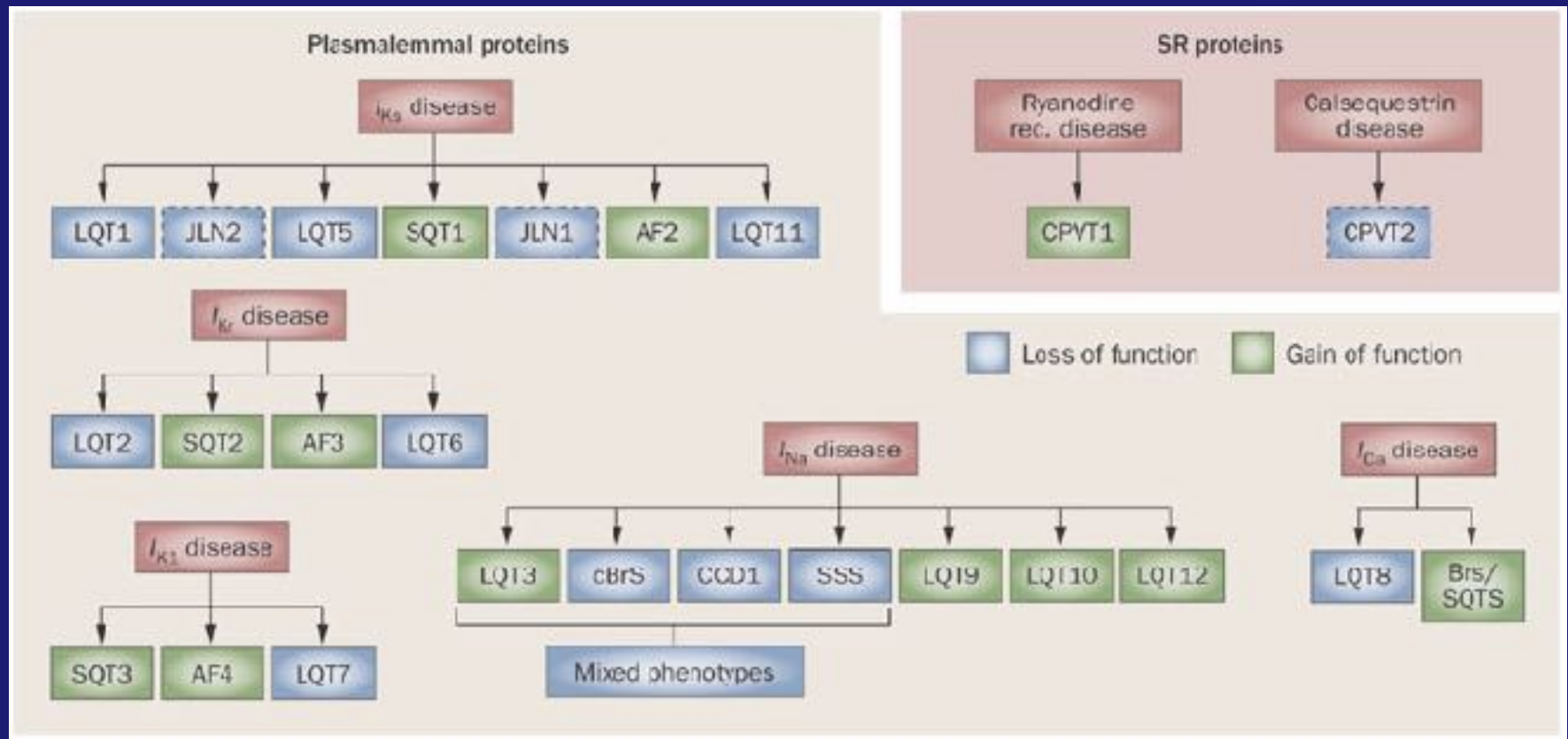


# Ionic Basis of the Action Potential in Mammalian Cardiomyocytes



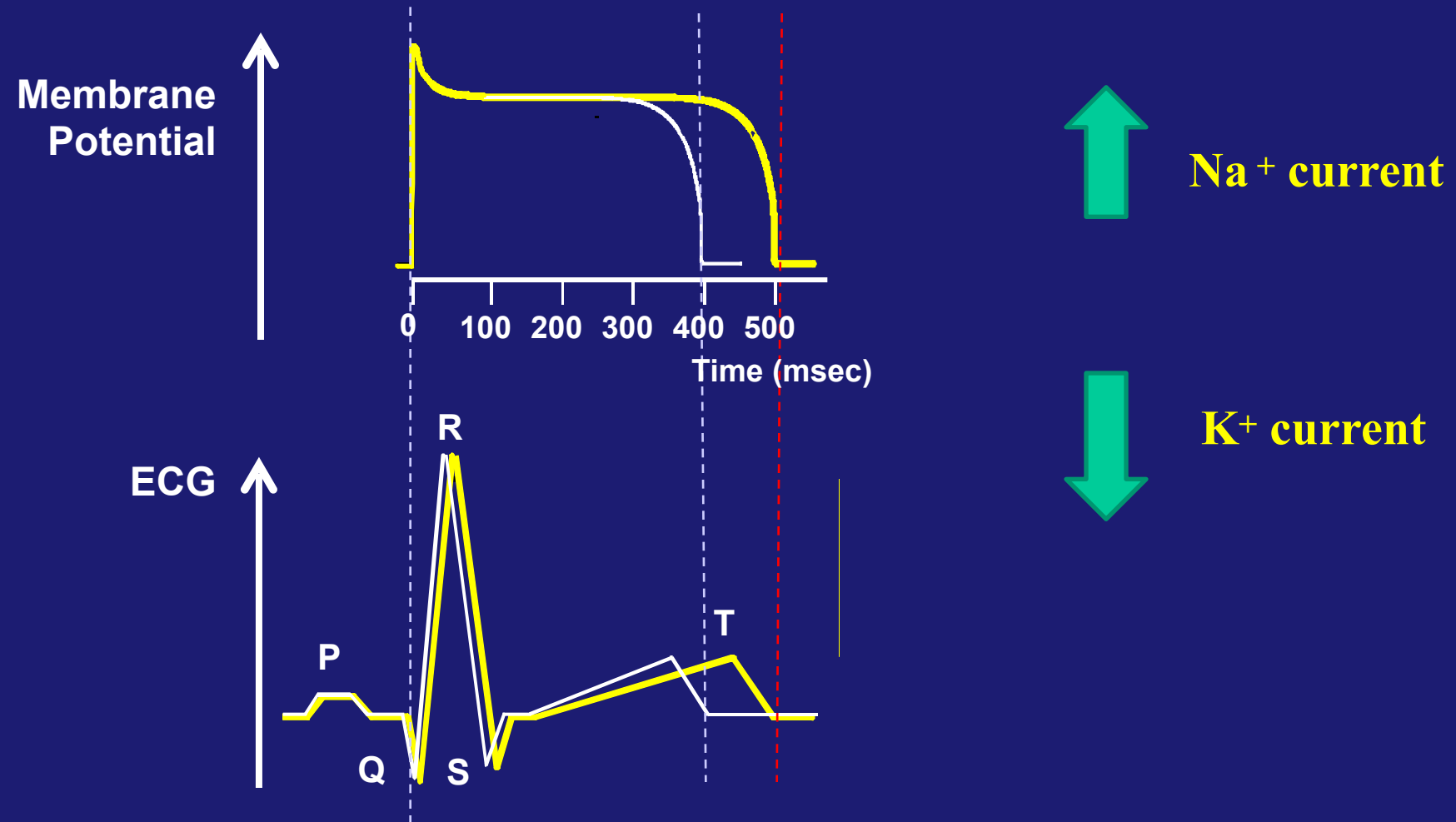


# Overlap in Gene Mutations



# Long QT Syndrome (LQTS)

# Long QT Syndrome



# Long QT Syndrome

- Congenital LQTS: 1 in 2500 to 10,000 in the general population
- SCD is relatively common (3000 to 4000 annual sudden deaths in childhood in the United States)
- An untreated mortality rate of 50% in 10 years.
- The primary symptoms in patients with LQTS include palpitations, syncope, seizures, and cardiac arrest.
- Majority of patients are asymptomatic
  - One third present with syncope or malignant arrhythmia (most common being torsades)

# Long QT Syndrome

- Acquired *versus* inherited
- Mutations in seven genes have been identified thus far in patients with genetic LQTS
- Two defined patterns of inheritance:
  - Autosomal Dominant (Romano-Ward Syndrome): Includes LQT Syndrome 1 to 12.
  - Autosomal Recessive (Jervell Lang-Nielsen Syndrome): Associated with deafness. Thus far has only been described in LQT1 and LQT5 (i.e. the genes encoding for  $I_{ks}$  current).

# Long QT Syndrome

- Andersen-Tawil syndrome (LQT7)
  - rare autosomal dominant ;
  - episodes of paralysis,
  - ventricular arrhythmias, and dysmorphic features ;
  - mutations in the gene KCNJ2, located on chromosome 17q23, which encodes the inward rectifying potassium channel Kir2.1
  - Characteristic T-U wave morphologies have also been identified in patients with Andersen syndrome:
    - Prolonged terminal T wave downslope
    - Biphasic U waves in limb leads
    - Wide T-U junction (in contrast to bifid T waves in LQT2)
    - Enlarged U waves

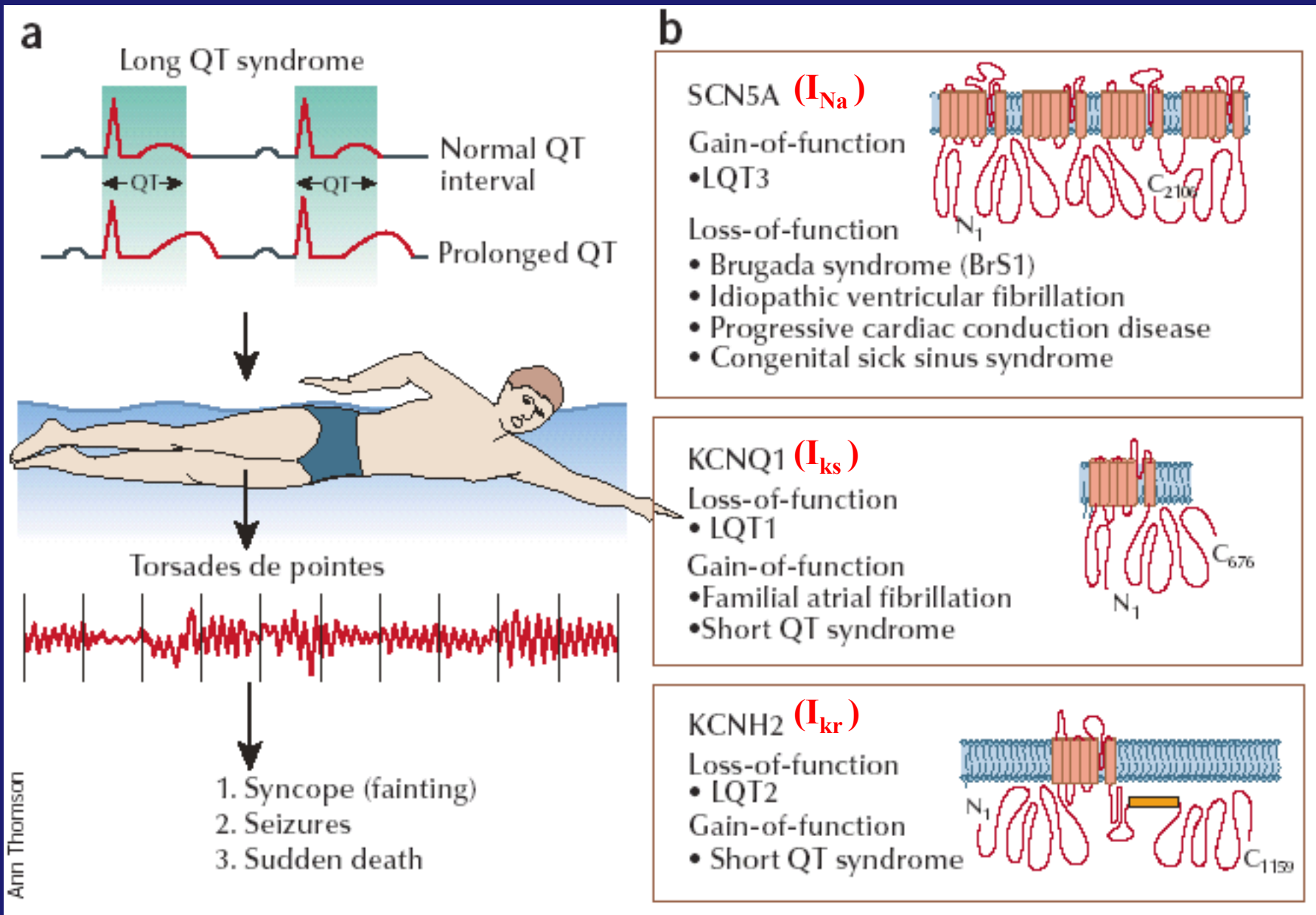
# Inherited LQTS

Locus name	Chromosomal locus	Gene symbol	Protein (symbol)	Current	Action potential	In vitro characterization	Gene-specific therapy*	Clinical syndrome = heterozygous mutation	Clinical syndrome = homozygous mutation
LQT1	11p15.5	KCNQ1	I <sub>Ks</sub> potassium channel $\alpha$ -subunit (Kv11.1)	↓ IKs	Delayed phase 3	Dominant negative suppression, trafficking defect, abnormal gating, reduced response to beta-AT signal	Beta-blockers*, potassium channel openers*	R-A	± L-H
LQT2	1q33-q38	KCNH2	I <sub>Kr</sub> potassium channel $\alpha$ -subunit (HERG)	↓ IKr	Delayed phase 3	Dominant negative suppression, trafficking defect, abnormal gating	Beta-blockers*, potassium supplement*, potassium channel openers*, mexiletine	R-A	NR
LQT3	3p21	SCN5A	Cardiac sodium channel $\alpha$ -subunit (Nav 1.5)	↑ INa	Prolonged phase 2	Abnormal gating: sustained current, slower inactivation, faster recovery, increased window current	Sodium channel blockers (mexiletine)*	R-A	NR
LQT4	4q25-q27	ANKK2	Ankyrin B (ANKB)	↓ Noct, Na/K ATPase, InsP3		Loss of expression and mislocalization	None proposed	R-A	NR
LQT5	21q22.1-q22.2	KCNK1	I <sub>Kd</sub> potassium channel $\alpha$ -subunit (MinK)	↓ IKd	Delayed phase 3	Dominant negative suppression, abnormal gating, reduced response to beta-AT signal	Beta-blockers, potassium supplement, potassium channel openers	R-A	± L-H
LQT6	21p13.1-q22.2	KCNK2	I <sub>K1</sub> potassium channel $\beta$ -subunit (IRX1)	↓ IK1	Delayed phase 3	Reduced current density and abnormal channel gating	Beta-blockers, potassium supplement, potassium channel openers, flecainide and thapsigargin	R-A	NR
LQT7/Neckless	17q23.1-q24.2	KCNK2	I <sub>K1</sub> potassium channel (IRX2.1)	↓ IK1	Delayed phase 3	Dominant negative suppression, nonfunctional channels, trafficking defect, abnormal gating	None proposed	H-P	NR
LQT8/Elmthy	12p13.3	CACOPHONY	Voltage-gated calcium channel, Cav1.2	↑ ICa	Delayed phase 3	Loss of inactivation	Calcium channel blockers*	R-A	None
LQT9	5p25	CACOPHONY	Caveolin-3	↑ INa		Increased late INa	Sodium channel blockers (mexiletine)		
LQT10	11q23	SCN5B	Cardiac sodium channel $\alpha$ -subunit 4	↑ INa	Prolong the action potential plateau	Increased late INa	Sodium channel blockers (mexiletine)	R-A and 2,3-ArB	
LQT11	1q21-q22	ANKK2	A-kinase anchoring protein	↓ IKs		Reduced phosphorylation of the IKs channel	Beta-blockers		
LQT12	20q11.2	SNCA	Syntaxin	↑ INa		Increased late INa	Sodium channel blockers (mexiletine)		

40-55%

35-45%

8-10%





# LQT: Triggers of Arrhythmia

- Genotype-phenotype correlations have been performed:
  - LQT1: Events related to swimming (occurring either immediately after diving into water or during recreational or competitive swimming activities)
  - LQT2: Events triggered by auditory stimuli, such as an alarm clock or telephone ringing
  - LQT3: At highest risk of events when at rest or asleep



# LQT: Prognosis

- The clinical course of LQTS is influenced by the specific analysis from the International Registry:
  - High risk ( $\geq 50$  percent) – patients with  $QT_c \geq 0.50$  sec who have LQT1 or LQT2 or (if male) LQT3
  - Intermediate risk (30 to 49 percent) – female patients with LQT3 and  $QT_c \geq 0.50$  sec or patients with  $QT_c < 0.50$  sec who have LQT3 or (if female) LQT2
  - Low risk ( $< 30$  percent) – patients with  $QT_c < 0.50$  sec who have LQT1 or (if male) LQT2

# LQT: Management

- The mortality rate in LQTS is reduced with earlier identification of affected patients and treatment:
  - beta blocker therapy,
  - sports restriction in some
  - avoidance of medications known to lengthen the QTc interval.
- High-risk patients including those with persistent symptoms despite beta blocker therapy may benefit from ICD implantation or left cardiac sympathetic denervation.
- Beta blockers are extremely protective in LQT1 patients, moderately protective in LQT2, and may not be sufficiently protective for those with LQT3. Consequently, targeting the pathologic, LQT3-associated late sodium current with agents such as mexiletine, flecainide, or ranolazine may represent a gene-specific therapeutic option for LQT3.

# Short QT syndrome

# Short QT syndrome

- Newly described in 2000
- Although not properly defined,  $QT < 330$  ms should raise high suspicion
- Three gene mutations identified so far:  $KCNH2$  ( $I_{kr}$ ),  $KCNQ1$  ( $I_{ks}$ ),  $KCNJ2$  ( $I_{ki}$ ).
- Clinically episodes of syncope, atrial fibrillation and/or life-threatening cardiac arrhythmias.
- Tx: ICD. Research underway with class III antiarrhythmics that prolong QT (eg. Quinidine).

# Brugada Syndrome

# Brugada Syndrome

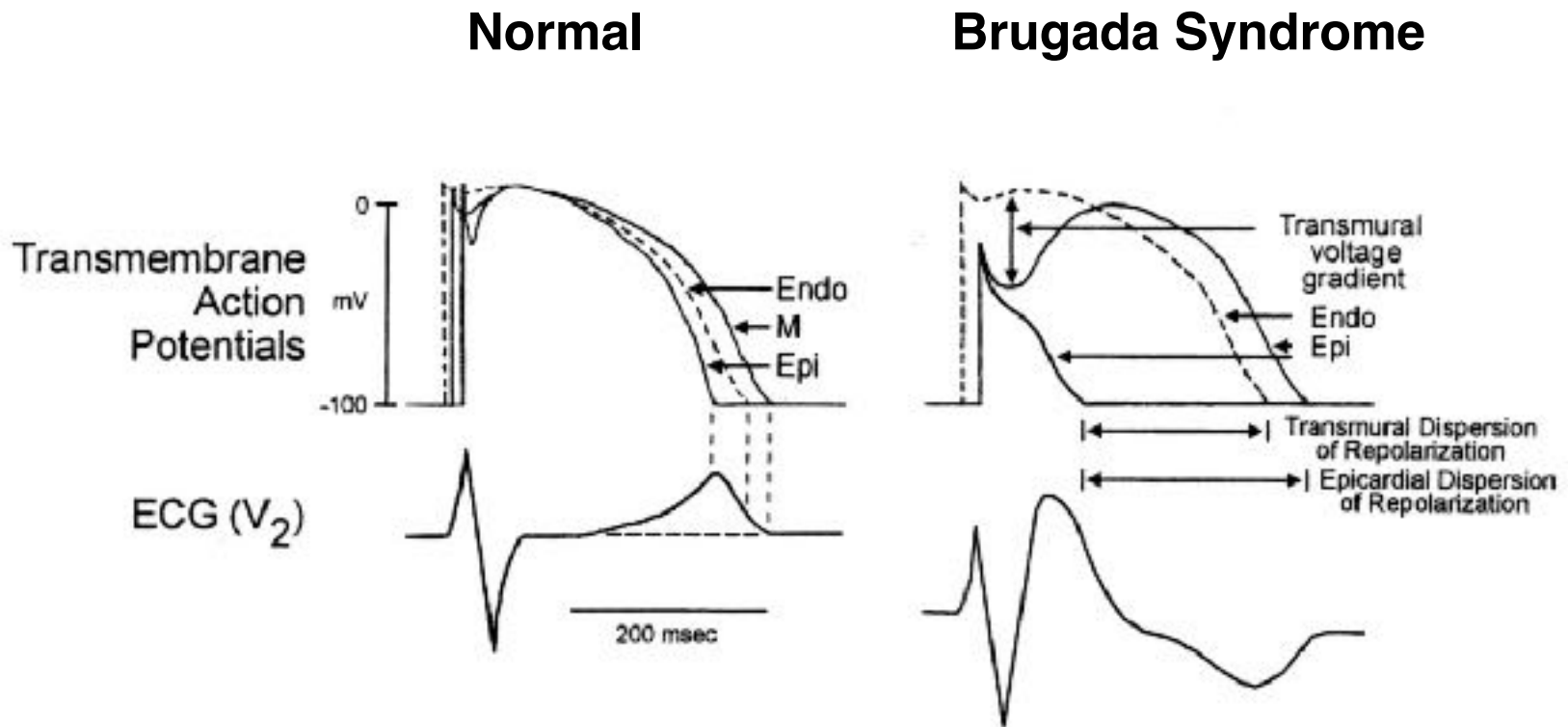
- Gene mutations of SCN5A (sodium channel), resulting in a loss of sodium channel function (either decreased expression or acceleration of inactivation) has been found in 10 to 30 percent of patients with Brugada syndrome.
  - In contrast, patients with congenital LQTS have SCN5A mutations that results in a gain in sodium channel function.
- Autosomal dominant transmission with incomplete penetrance
- A distinct clinical syndrome with syncope episodes and sudden cardiac death (fast polymorphic VT)

# Brugada Syndrome

- This syndrome is estimated to be responsible for 20-50% of all sudden death in patients with an apparently normal heart.
- Prevalence varies from 5-50:10,000; largely depending on geographic location (endemic in southeast Asia).
- SUDS (Sudden Unexpected Death Syndrome) in Southeast Asia is a form of Brugada syndrome; most common cause of death in young males in Thailand.
- Male: Female 10:1



# Loss of Normal Heterogeneity with Increased Dispersion of Repolarization



The right ventricle is most affected in Brugada syndrome, and particularly (but not specifically) the right ventricular outflow tract.

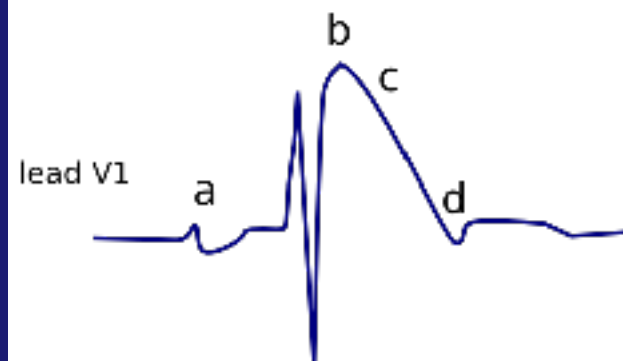
12-lead electrocardiogram (ECG) from a patient with the Brugada syndrome shows downsloping ST elevation



ST segment elevation and T wave inversion in the right precordial leads V1 and V2 (arrows); the QRS is normal. The widened S wave in left lateral leads that is characteristic of right bundle branch block is absent.

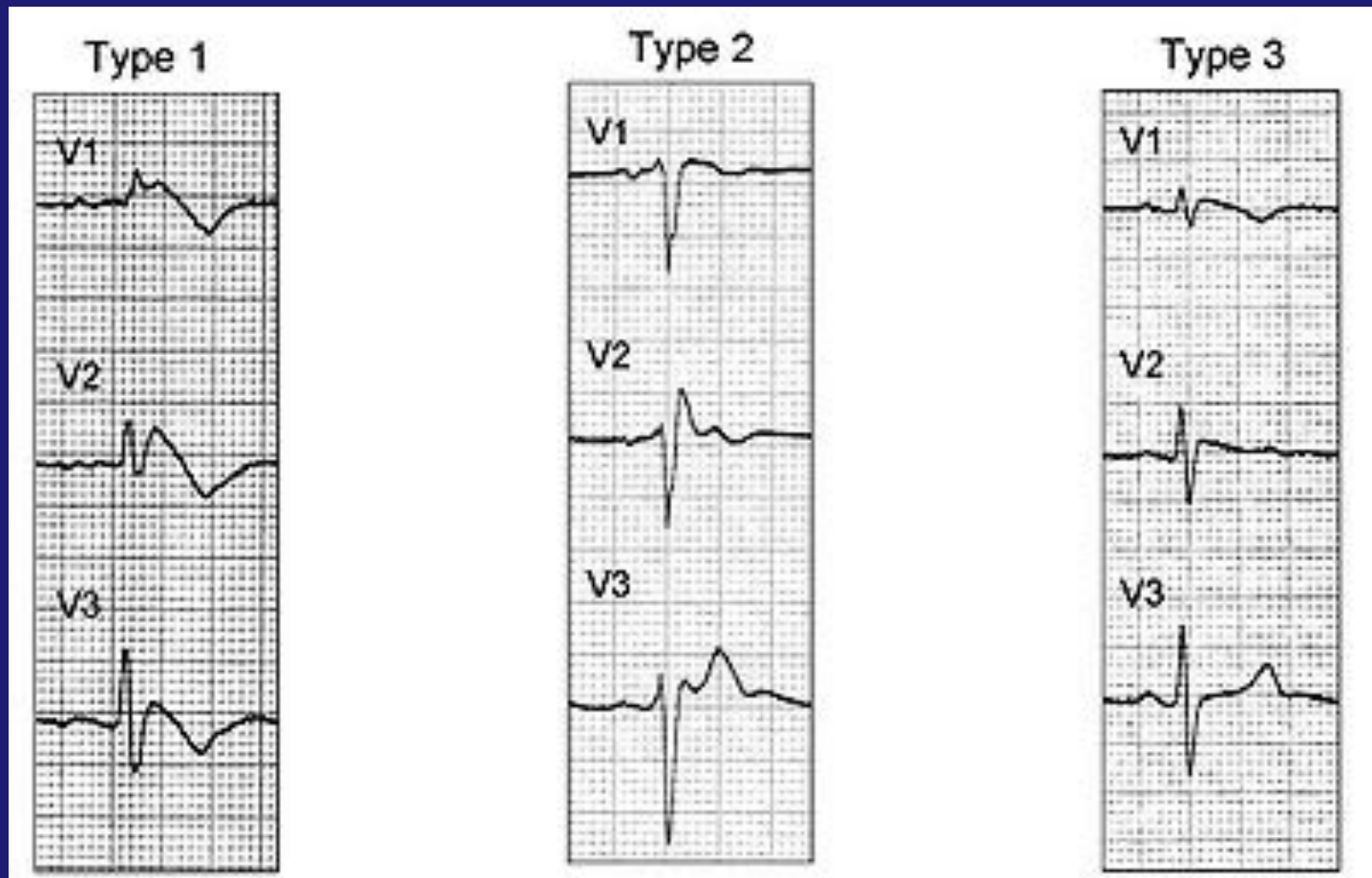
*Courtesy of Dr Rory Chikara, University of Chicago.*

RBBB pattern and >2mm ST segment elevation in V1-V3 in the absence of ischemia, electrolyte imbalance, and structural heart disease



ECG characteristics in Brugada Syndrome  
a. Broad P wave with some PQ prolongation  
b. J point elevation  
c. Coved type ST segment elevation  
d. Inverted T wave

# Brugada Syndrome: ECG



# Brugada Syndrome: Factors / Drugs that Enhance ECG Pattern

- Na<sup>+</sup> channel blockers
- alpha agonists, vagotonic agents, beta blockers
- fever
- alcohol, cocaine
- severe ischemia
- tricyclic antidepressants, antihistaminics

# Brugada Syndrome: Diagnostic Criteria

- **Major criteria:**

- 1. presence of ECG marker (Type 1) in structurally normal hearts
- 2. appearance of ECG marker after administration of Na<sup>+</sup> channel blockers (Type 2&3)

- **Minor criteria:**

- 1. family history of sudden cardiac death
- 2. syncope of unknown origin
- 3. documented ventricular tachycardia/fibrillation
- 4. genetic mutation of ion channels

## Brugada Syndrome

## Long QT Syndrome

### Intrinsic Heterogeneity

↑ Net outward Current  
Preferential Abbreviation  
of Epicardial APD

↓ Net outward Current  
Preferential prolongation  
of M Cell APD

↑ Dispersion of Repolarization  
(Epicardial and Transmural)  
QT interval ↔  
ST Segment Elevation

↑ Dispersion of Repolarization  
(Transmural)  
QT Interval Prolonged  
ST Segment ↔

Phase 2 reentrant  
Extrasystole

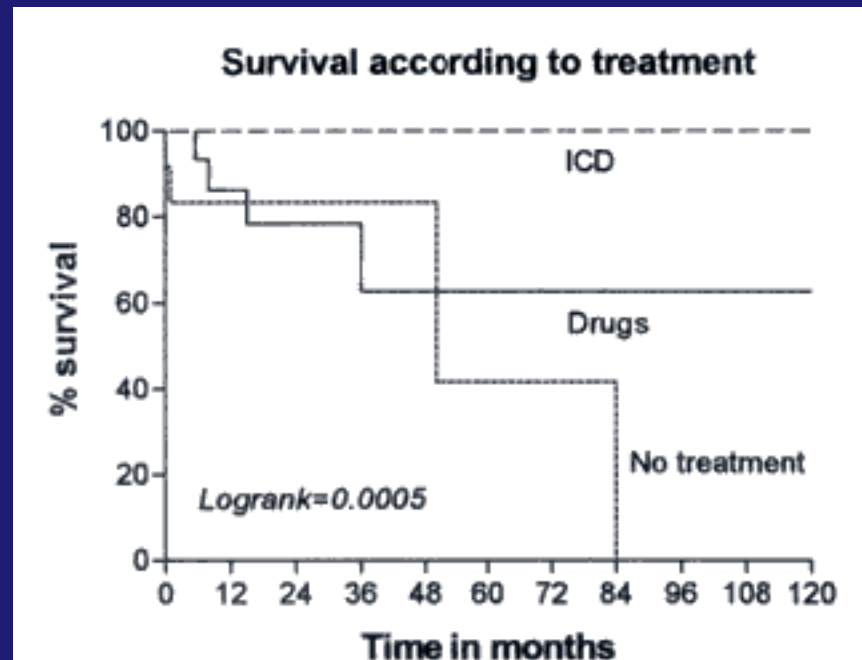
EAD-induced  
Extrasystole

Rapid Polymorphic VT/VF  
(reentry)

Polymorphic VT  
Torsade de Pointes  
(reentry)

# Brugada: Management

- Although pharmacologic therapy has been tried, the only therapy with proven efficacy in preventing sudden death is an implantable cardioverter-defibrillator (ICD).

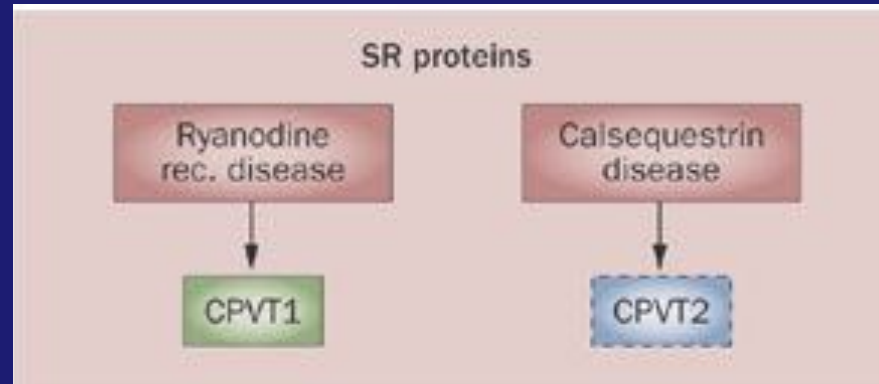




# Catecholaminergic Polymorphic Ventricular Tachycardia (CPVT)



# CPVT

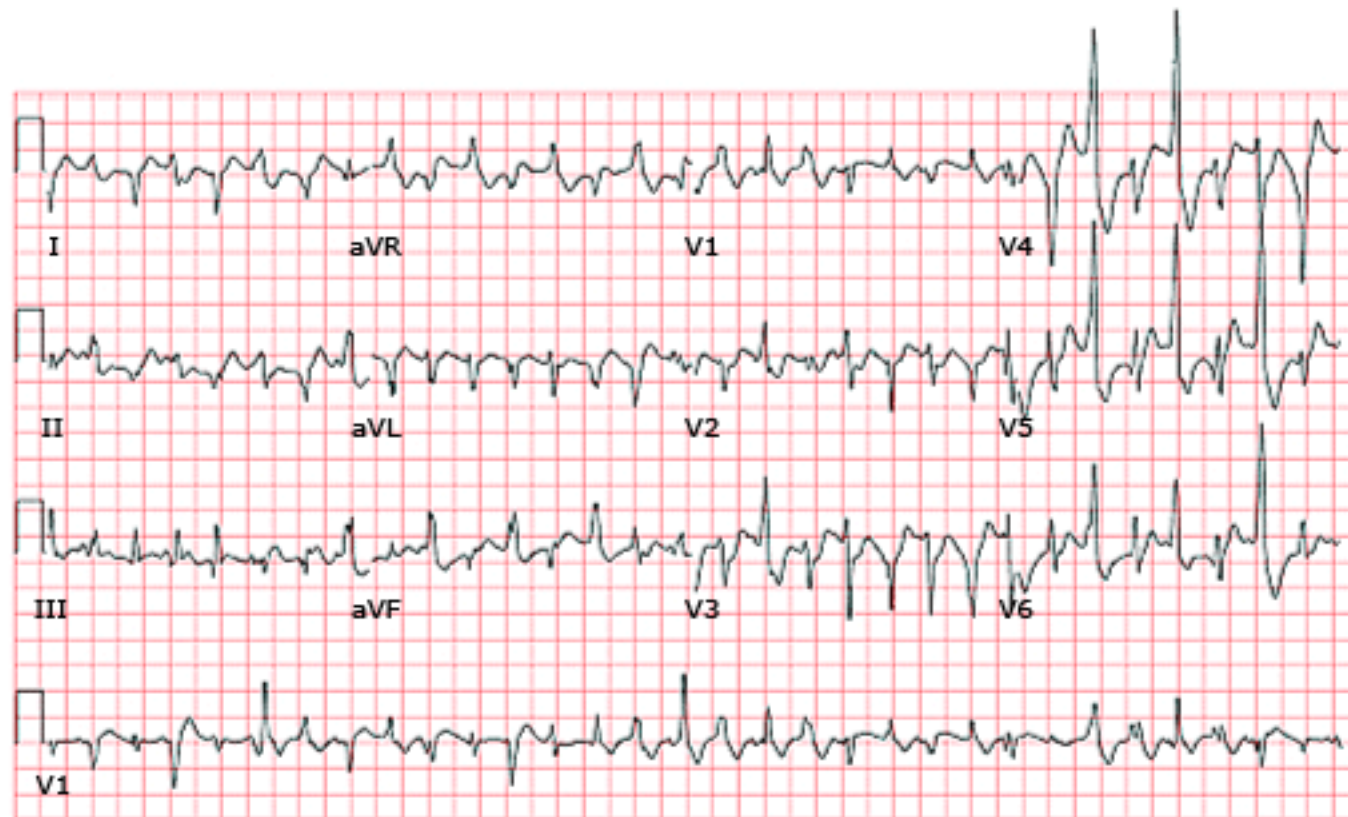


- Ryanodine receptors (RyR2 gene)
  - Responsible for release of calcium from the sarcoplasmic reticulum and are activated by incoming calcium; therefore they are activated Ca channels.
  - Autosomal Dominant
- Calsequestrin 2 gene
  - Encodes a major calcium reservoir protein within the sarcoplasmic reticulum.
  - Autosomal recessive

# CPVT

- At risk for VT (specifically bidirectional VT), ventricular fibrillation, and sudden death, especially in association with stress or exercise.
- At rest, the ECG typically demonstrates frequent premature ventricular contractions (PVC) and nonsustained polymorphic VT.
  - The QT interval is normal, which distinguishes it from LQTS, and there are no ST segment changes differentiating it from Brugada syndrome.
- At present, therapies include beta blockers, flecainide, and ICD or left cardiac sympathetic denervation for high risk patients and those with symptoms despite beta blocker therapy.
- Avoid competitive sports

## 12-lead ECG in child with catecholaminergic polymorphic ventricular tachycardia (CPVT)



25mm/s 10mm/mV 150Hz MUSE 7.0.0 12SL 250 CID: 1

EID:3087 EDT: 10:22 07-MAR-2002 ORDER:

This is the ECG of a five-year-old with recurrent exertional syncope. The ECG demonstrates bidirectional VT. This child underwent genetic testing and was positive for the RyR2 mutation.

# Idiopathic Ventricular Fibrillation

- Absence of identifying structural or genetic abnormalities to explain the VF or the out-of-hospital cardiac arrest
- May account for as much as 10% of sudden deaths, especially in the young.
- About 30% of IVF-labeled individuals will have recurrent episodes of VF.
- Like BrS, loss of function SCN5A mutations have been identified.
  - Identified mutations in other arrhythmia susceptibility genes, such as *ANKK*, which encodes for ankyrin-B, and *RYR2*, which encodes for the cardiac ryanodine receptor.

# Idiopathic V-fib

- J-point elevation (1 mm above baseline) on inferolateral electrocardiographic leads (so-called early repolarization)
  - was significantly overrepresented (31%)
  - was greater in magnitude in subjects who experienced cardiac arrest caused by IVF compared with age-, gender-, race-, and level of physical activity–matched controls.
- These patients with early repolarization were more often males and had a personal history of syncope or cardiac arrest during sleep

# Progressive Cardiac Conduction Defect

- Lev-Lenègre disease
- Progressive (age-related) alteration of impulse propagation through the His-Purkinje system, with right or left bundle branch block and widening of the QRS complex, leading to complete atrioventricular (AV) block, syncope, and occasionally sudden death.
- Over 30 PCCD-associated mutations in *SCN5A* (*loss of function*), *autosomal dominant*.
- PCCD is the prevailing phenotype in BrS-associated *SCN5A* mutation carriers, where the penetrance of conduction defects was 76%.

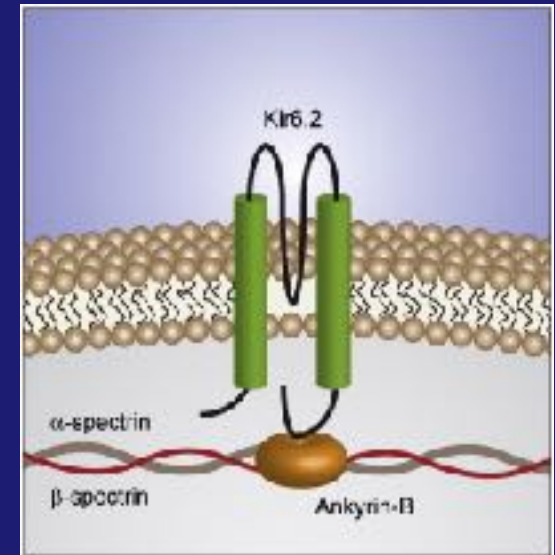
# Sick Sinus Syndrome

- Inappropriate sinus bradycardia, sinus arrest, atrial standstill, tachycardia-bradycardia syndrome, or chronotropic incompetence
- Acquired: Commonly occurs in older adults (1 in 600 cardiac patients older than 65 years) with acquired cardiac conditions.
- Idiopathic: No identifiable cardiac anomalies; can occur at any age, including in utero.
  - Additionally, familial forms of idiopathic SND consistent with autosomal dominant inheritance with reduced penetrance and recessive forms with complete penetrance have been reported.
- So far implicated three genes—*SCN5A*, *HCN4* (encodes the so-called  $I_f$  or pacemaker current and plays a key role in automaticity of the sinus node), and *ANKB*.

# Ankyrin-B Syndrome

*ANK2* gene → Ankyrin-B protein

Involved in anchoring the Na<sup>+</sup>,K<sup>+</sup>-ATPase, Na<sup>+</sup>/Ca<sup>2+</sup> exchanger, and InsP3 receptor to specialized microdomains in the cardiomyocyte transverse tubules.



- Loss of function mutations of *ANK2* were shown originally to cause a dominantly inherited cardiac arrhythmia with an increased risk for SCD associated with a prolonged QT interval
- LQT4 → more correctly renamed SSS with bradycardia, or the ankyrin-B syndrome.



# References

- Braunwald's Heart Disease Textbook, 9<sup>th</sup> Edition
- Hurst's The Heart Textbook, 13<sup>th</sup> Edition

A collection of orange and white Tennessee Volunteers merchandise. In the background, a large white bag with a large orange 'T' and a checkered pattern. In the foreground, a wooden sign with 'VOLS' written vertically, two clear plastic containers with orange lids and 'WAX' labels, a wooden sign with 'KT' written vertically, and a small orange and white checkered bag with a 'T' on it. The items are arranged on a white surface with a repeating pattern of the Tennessee Volunteers logo.