

The Application of Molecular Genetics in Sudden Death and Near Death – the Social and Medico-legal Impacts

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SUDDEN DEATH AND CARDIAC DISEASES

Definition

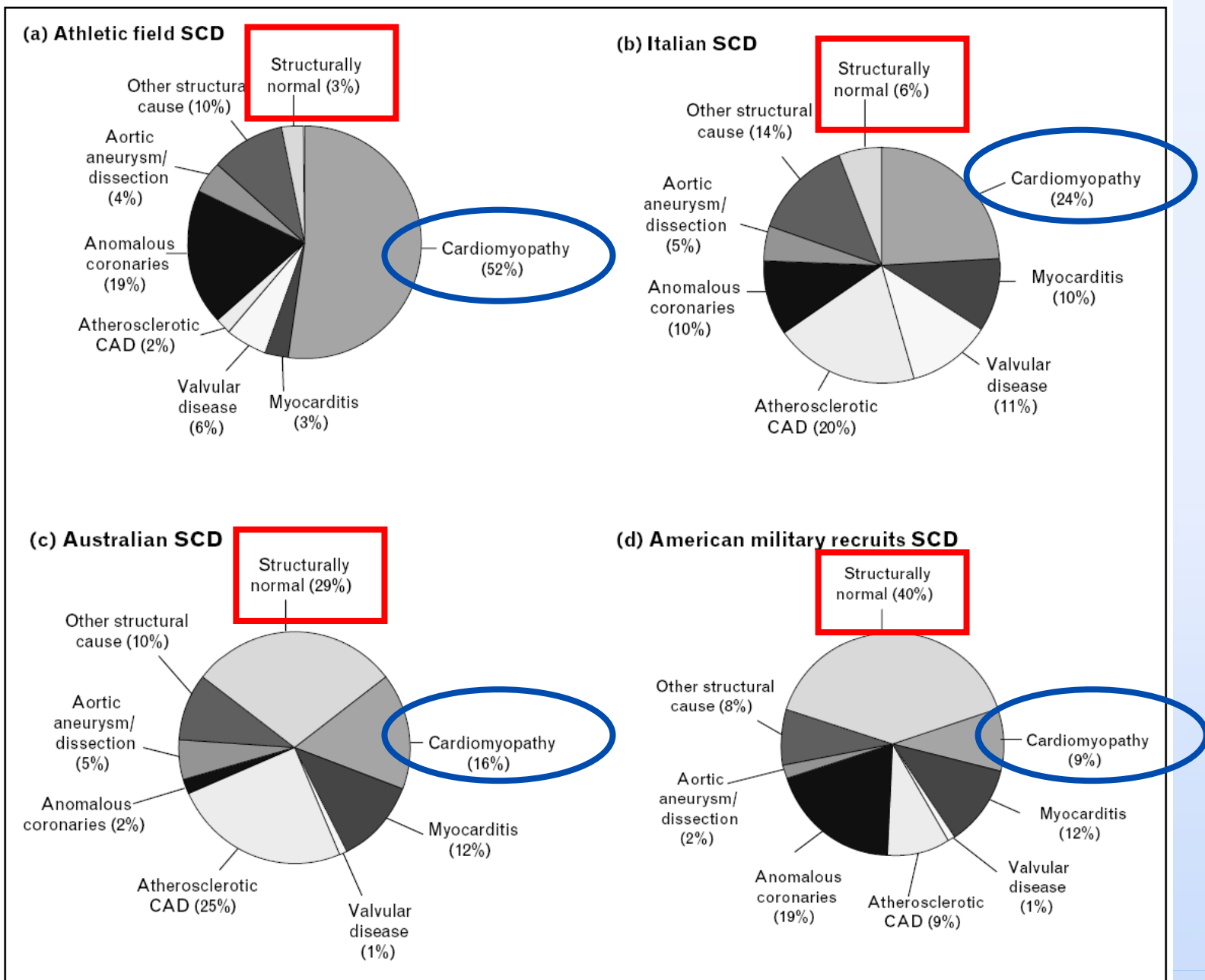
* Sudden death -

- * “an unexpected natural death within a short time period, generally 1 hour or less from the onset of symptoms” or
- * “a non-witnessed death discovered within 24 hours in someone without prior symptoms or any prior condition that would appear fatal”.

Sudden Death

- * Incidence and causes are age-dependent.
- * Coronary artery disease in adults > 35y.
- * Accounts for 5% of all deaths in children and adolescents with an incidence of 1.5 – 8 per 100,000 patient-year.

Figure 1 Causes of sudden cardiac death in young people



Heritable causes of sudden cardiac death

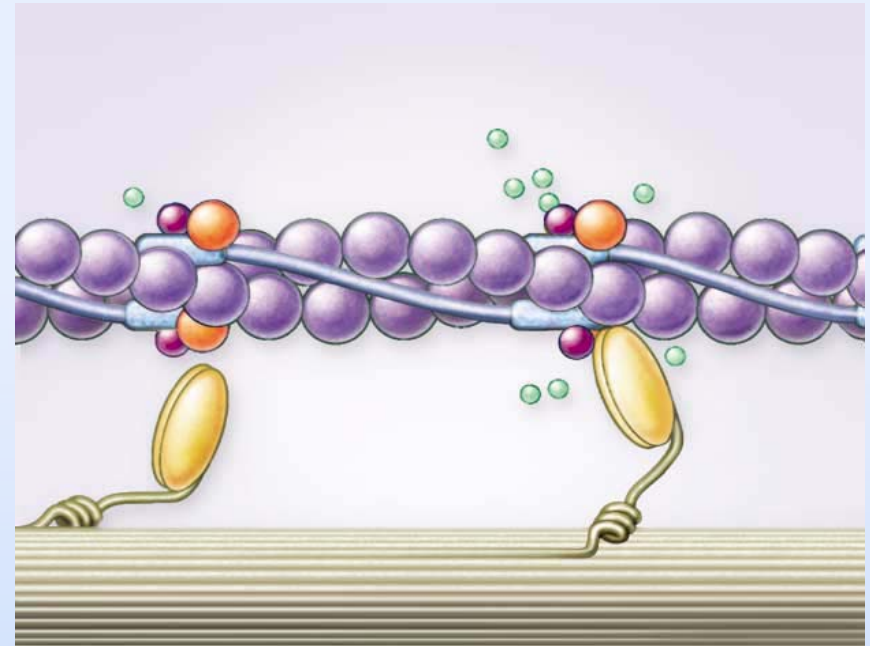
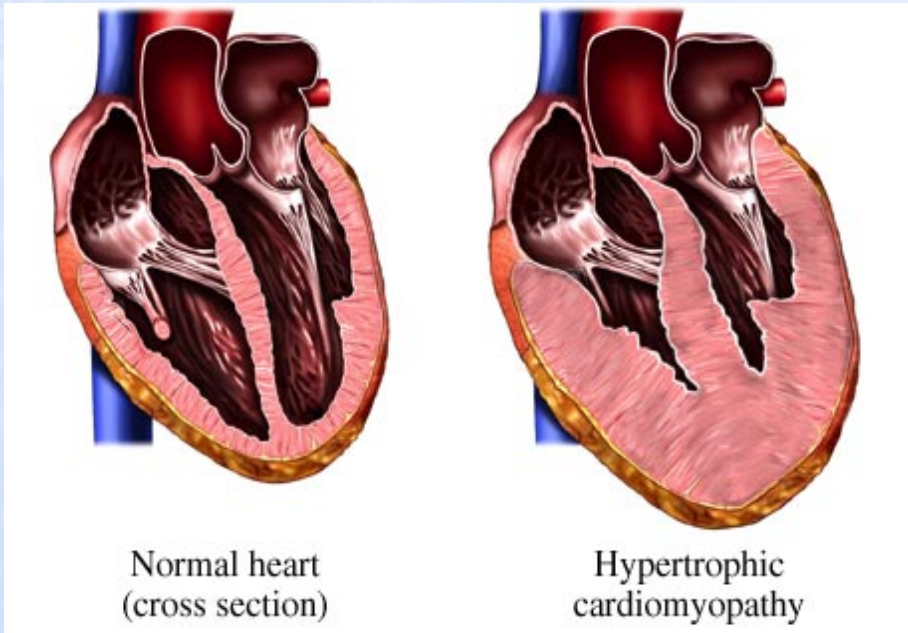
Structural heart defects

- ✿ Hypertrophic cardiomyopathies (HCM)
- ✿ Arrhythmogenic right ventricular cardiomyopathy (ARVC)

Cardiac electrical disorders

- ✿ Long QT syndromes (LQTS)
- ✿ Catecholaminergic polymorphic ventricular tachycardia (CPVT)
- ✿ Brugada syndrome

Hypertrophic Cardiomyopathy

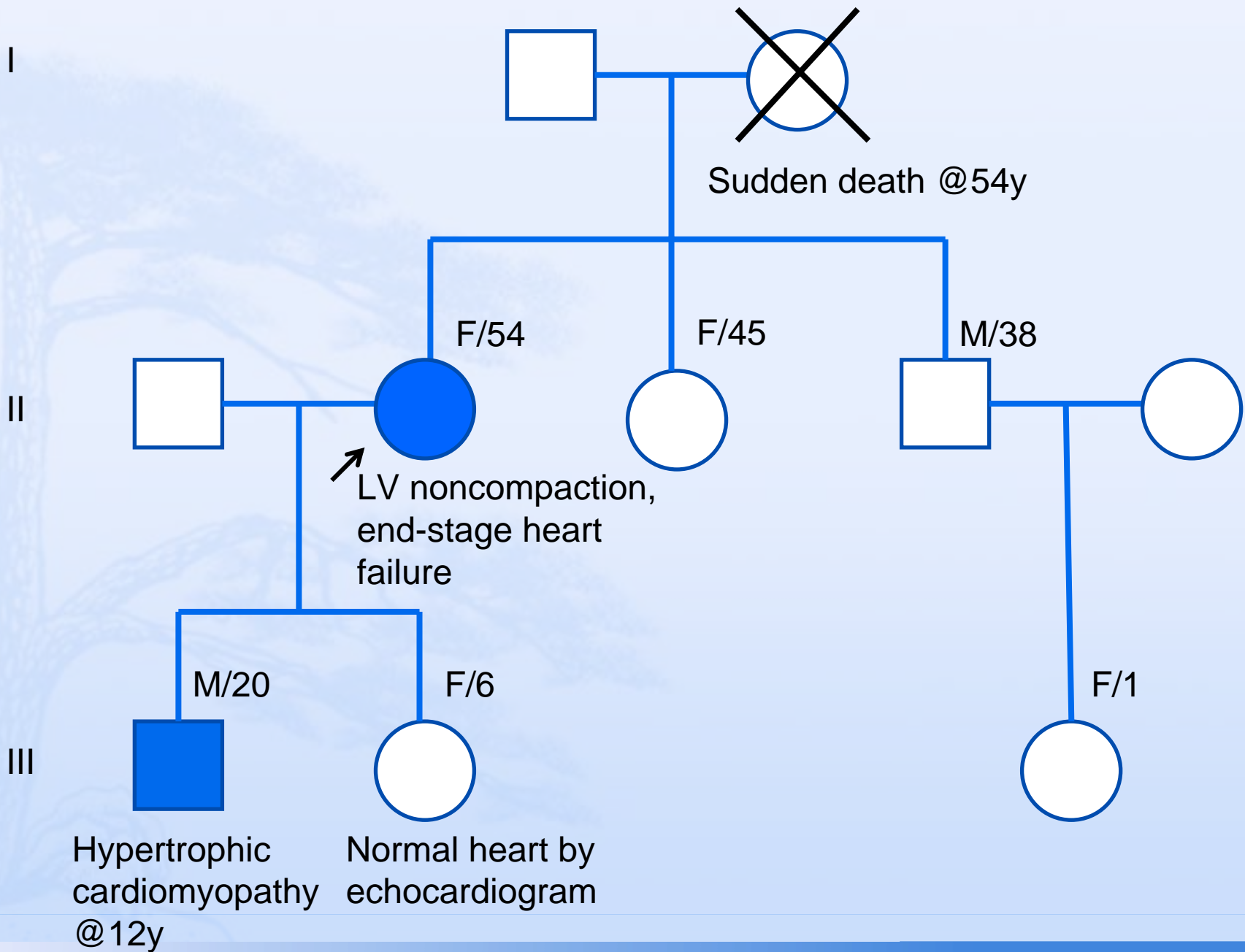


Sarcomere

HCM

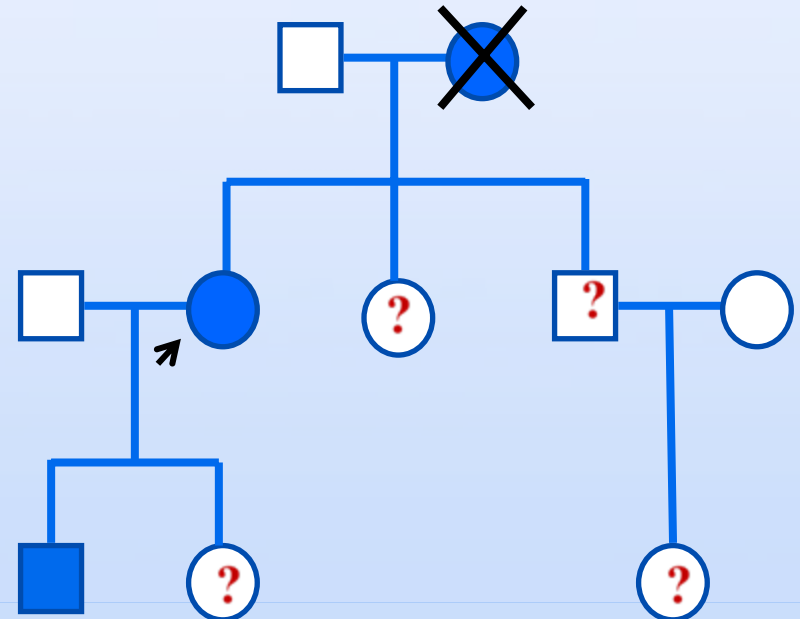
- ✿ Account for ~1/3 of sudden deaths in competitive athletes in the US .
- ✿ 55%-70% attributable to one of the 12 sarcomere genes.
- ✿ Autosomal dominant with age-related penetrance
- ✿ Existing evidence: diagnosis and family screening
- ✿ Little prognostic value, cannot predict age of onset and disease severity.

Locus Name	Gene Symbol	Protein Name	% of HCM Caused by Mutations in This Gene
CMH1	MYH7	Myosin heavy chain, cardiac muscle beta isoform	40%
CMH4	MYBPC3	Myosin-binding protein C, cardiac-type	40%
CMH2	TNNT2	Troponin T, cardiac muscle	5%
CMH7	TNNI3	Troponin I, cardiac muscle	5%
CMH3	TPM1	Tropomyosin 1 alpha chain	2%
CMH10	MYL2	Myosin regulatory light chain 2, ventricular/cardiac muscle isoform	Unknown
CMH8	MYL3	Myosin light polypeptide 3	1%
	ACTC1	Actin, alpha cardiac muscle 1	Unknown
	CSRP3	Cysteine and glycine-rich protein 3, muscle LIM protein	Unknown
CMH9	TTN	Titin	
	MYH6	Myosin heavy chain, cardiac muscle alpha isoform	
	TCAP	Telothonin	
Other genes implicated in HCM			
	TNNC1	Troponin C, slow skeletal and cardiac muscles	Unknown



This pedigree

- ✿ Familial disease
- ✿ Dominant inheritance
- ✿ Phenotypes: childhood-onset HCM, end-stage heart failure, (sudden death)
- ✿ At risk asymptomatic family members



Clinical vs Genetics Dx

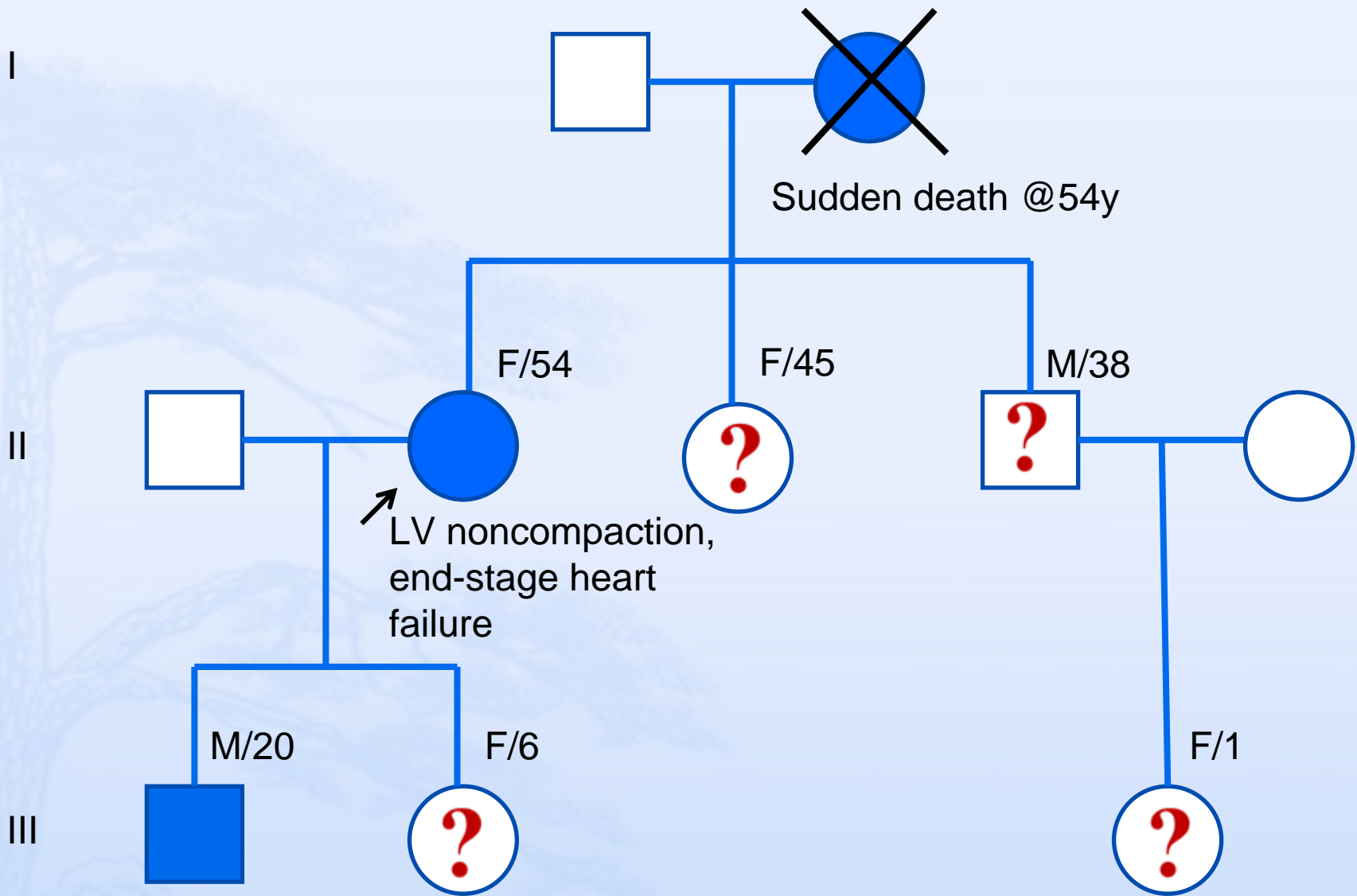
Clinical Screening

- ✿ History, physical exam, echocardiography, ECG, etc.
- ✿ Always indicated in symptomatic individuals regardless of age.
- ✿ **A normal baseline echocardiogram and ECG does not rule out HCM in asymptomatic relatives, particularly in children or young adults.**
- ✿ **Require screening and longitudinal FU throughout life.**

Genetics vs Clinical Dx

Genetic Diagnosis

- * Testing of at-risk asymptomatic relatives is possible if the disease-causing mutation in the proband is known.
- * Genetically heterogeneous
- * Mutation detection rate 55-70%
- * **A negative test result can provide reassurance that the person is not at risk of developing HCM and thus obviate unnecessary screening.**



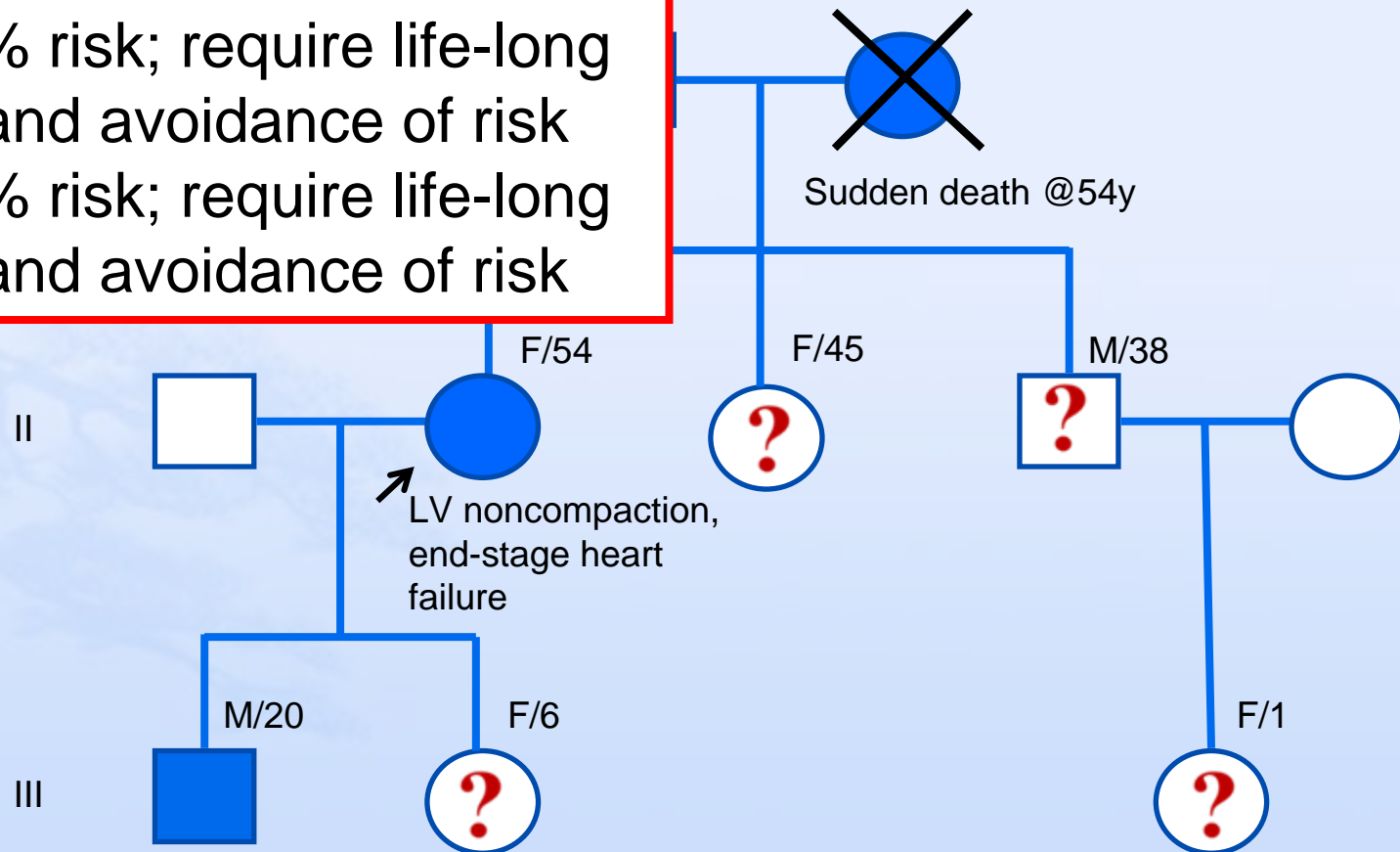
Hypertrophic cardiomyopathy @12y

Normal heart by echocardiogram

At risk asymptomatic relatives

Clinical screening

- III:1 – symptomatic
- II:3 and II:4 – 50% risk; require life-long follow-up and avoidance of risk
- III:2 – 50% risk; require life-long follow-up and avoidance of risk
- III:3 – 25% risk; require life-long follow-up and avoidance of risk



Hypertrophic cardiomyopathy @12y

Genetic Diagnosis

- ✿ Target 1: **MYH7**
- ✿ The beta heavy chain subunit of human muscle myosin
- ✿ Chromosome 14q12
- ✿ 38 coding exons
- ✿ ~300 known mutations throughout the gene, majority being missense / nonsense mutations.
- ✿ Account for ~40% familial HCM in Chinese



Sample collection



DNA Extraction



PCR



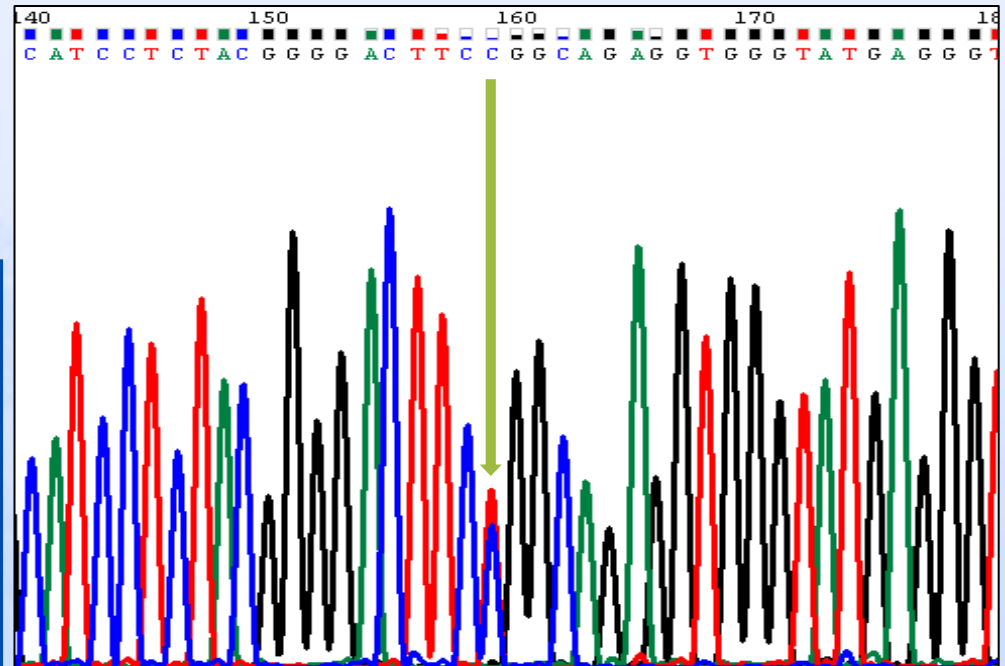
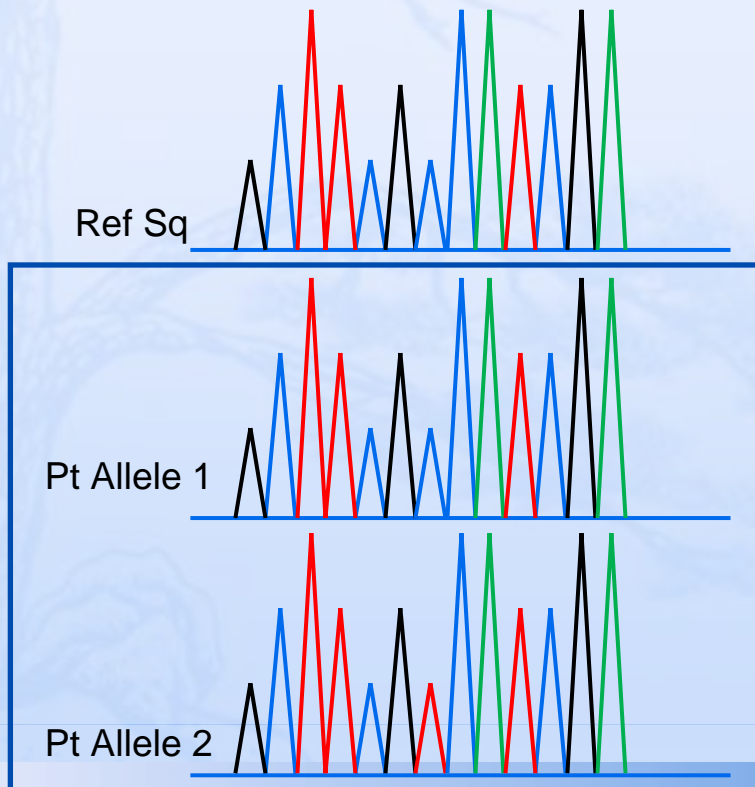
Sequencing Reaction

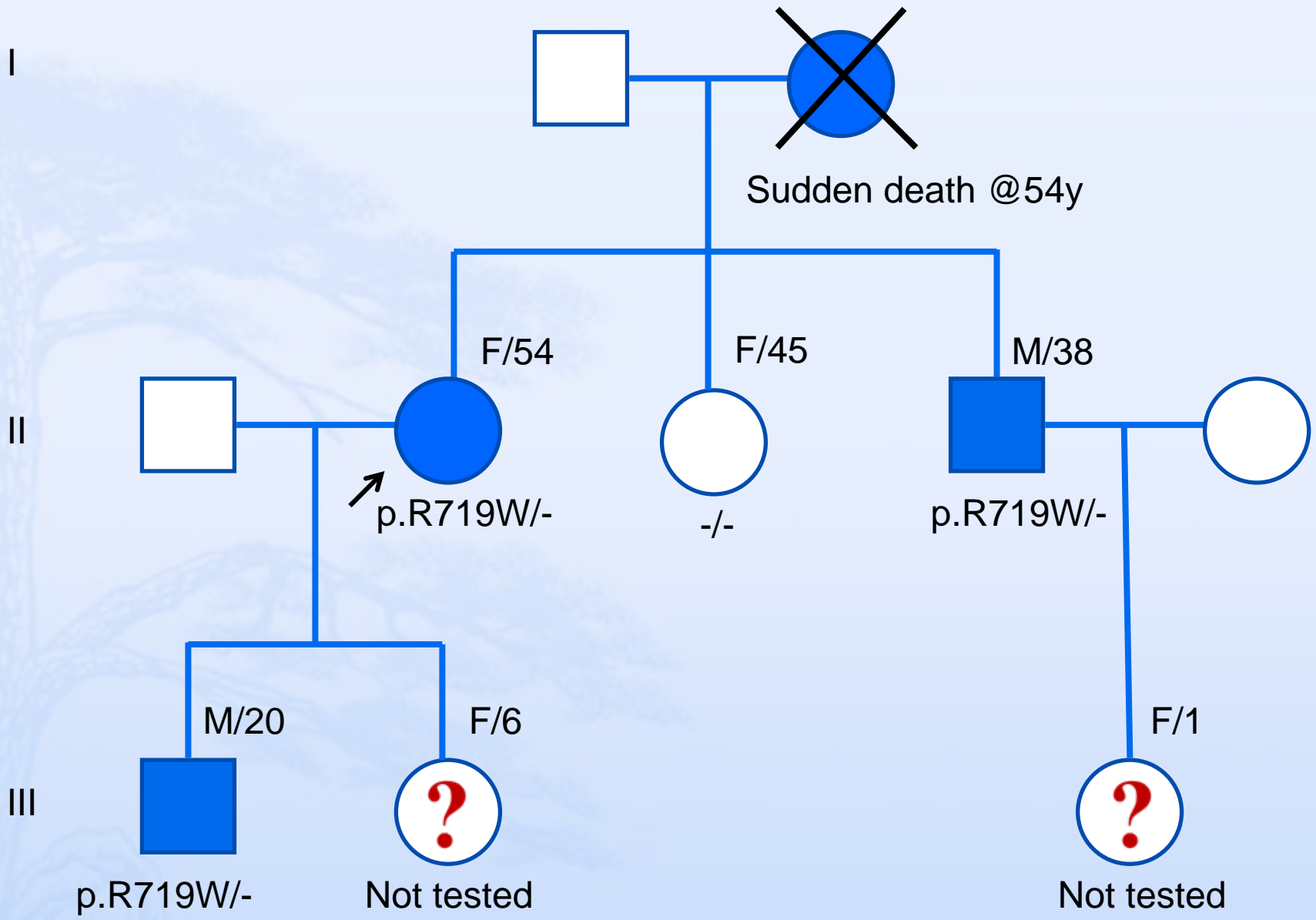


Sequencing

Sequencing results

- * c.2155C>T p.Arg719Trp in exon 19
- * First reported in 1994



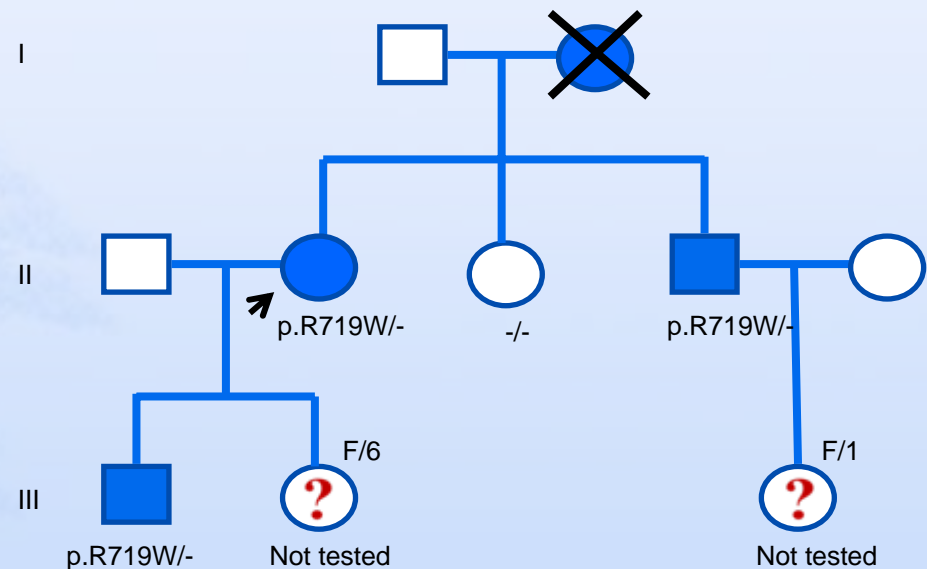


Clinical screening

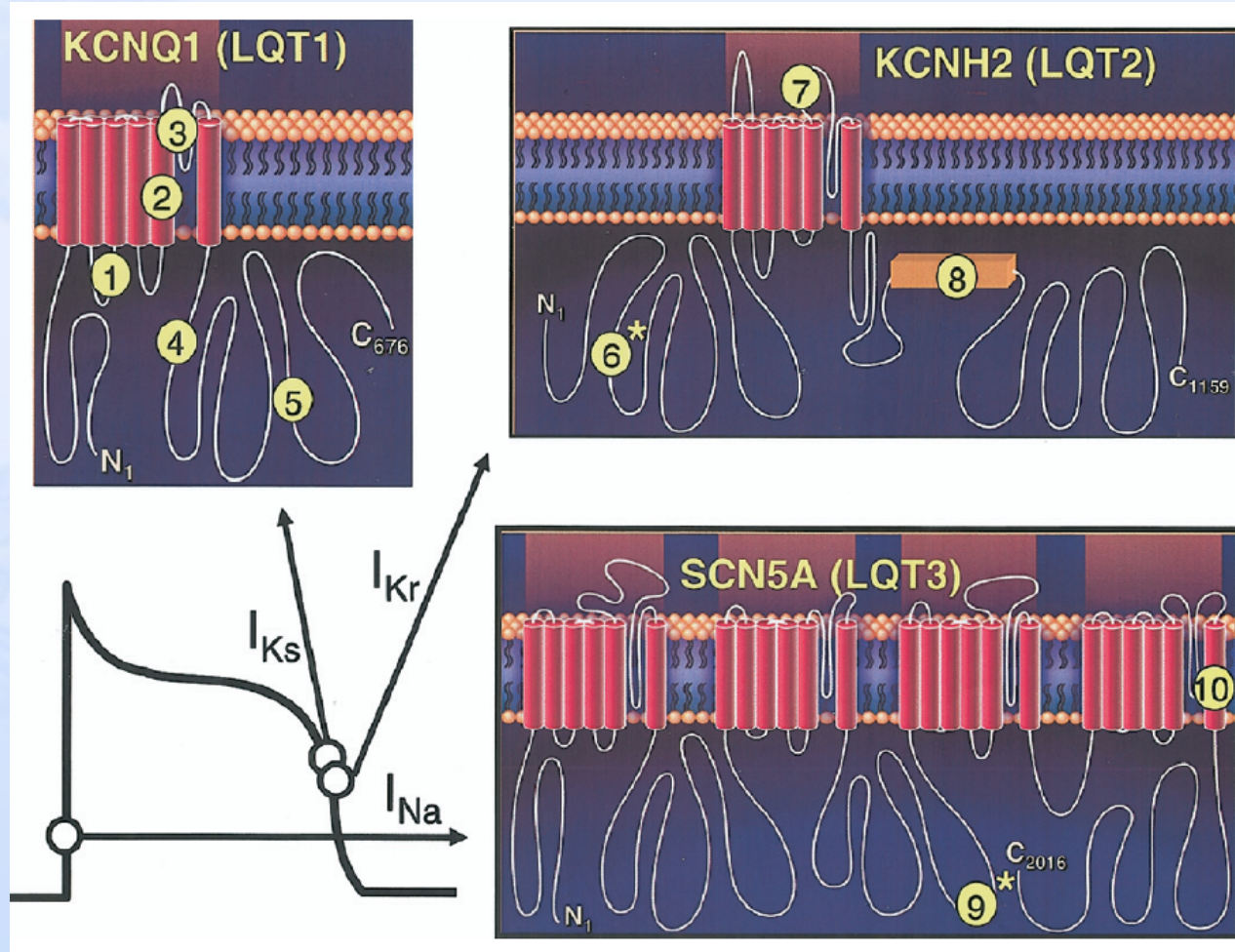
- III:1 – symptomatic
- II:3 – 50% risk
- II:4 – 50% risk
- III:2 – 50% risk
- III:3 – 25% risk

Genetic screening

- III:1 – same
- II:3 not at risk
- II:4 Dx confirmed
- III:2 – 50% risk
- III:3 – 50% risk



Cardiac Electrical Disorders

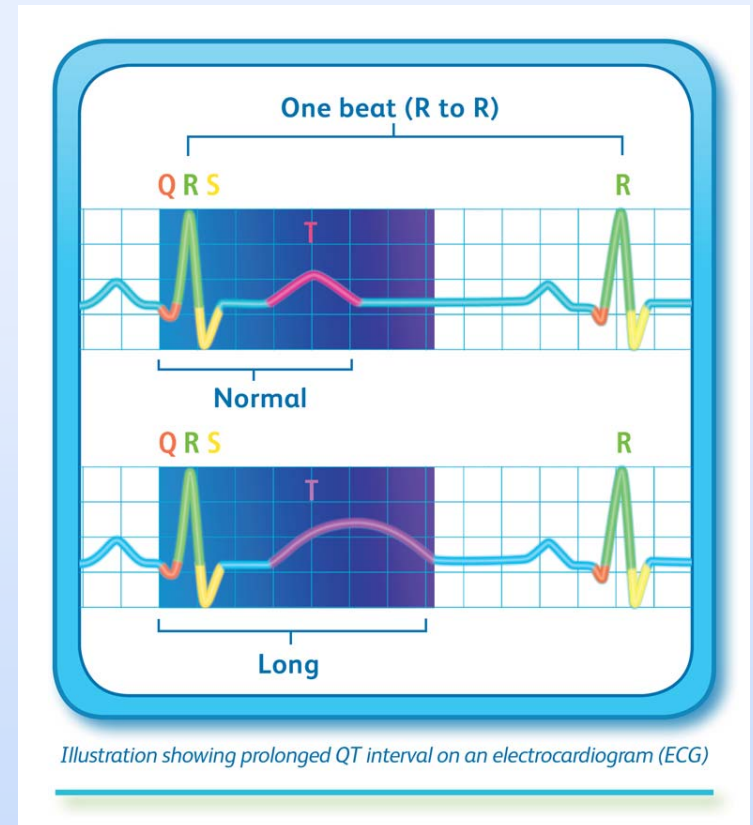


Cardiac electrical disorders

- * Cardiac channelopathies
- * No detectable structural abnormalities
- * Long QT syndromes (LQTS)
- * Catecholaminergic polymorphic ventricular tachycardia (CPVT)
- * Brugada syndrome (BrS)

Congenital LQTS

- * Autosomal dominant with reduced penetrance
- * Genetically heterogeneous
- * 75% attributable to
 - * KCNQ1 (LQT1)
 - * KCNH2 (LQT2)
 - * SCN5A (LQT3)
 - * KCNE1 (LQT5)
 - * KCNE2 (LQT6)



LQTS

	LQT1	LQT2	LQT3
Gene / Locus	KCNQ1 11p15.5	KCNH2 7q35-q36	SCN5A 3p21
Coding exons	16	15	27
Arrhythmogenic triggers	<ul style="list-style-type: none"> • Physical exertion • Swimming 	<ul style="list-style-type: none"> • Auditory stimuli (e.g. alarm clock) • Post-partum period 	<ul style="list-style-type: none"> • Sleeping
Response to β-blocker	++++	+++	+/-

Identification of asymptomatic LQTS carriers

- ✿ Clinical criteria are insensitive
 - ✿ A Schwartz score of ≥ 4 (i.e. a strong probability of LQTS) had a sensitivity of 38%.
 - ✿ Clinical assessment failed to identify 40% of the genetically affected family members of 310 genotyped probands.
- ✿ Patients with “concealed” LQTS have a 10% risk of cardiac events by age 40 years if left untreated.
- ✿ Early identification
 - ✿ Preventive life-style modifications (e.g. avoidance of strenuous exercise and extreme emotional stress)
 - ✿ Prophylactic treatment.
- ✿ Genetic test \Rightarrow gold standard in family screening

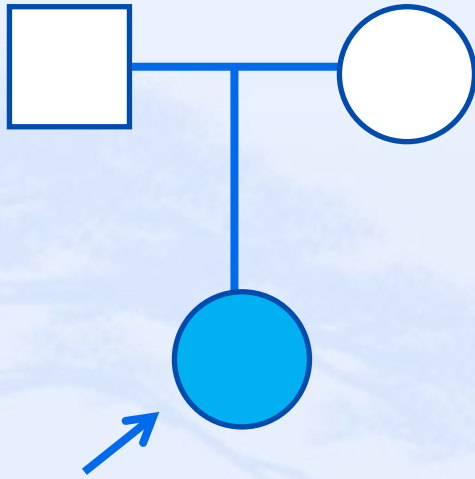
CPVT

- * Catecholaminergic polymorphic ventricular tachycardia
- * Syncope and sudden death during physical exertion or emotion (Catecholamine-induced bi-directional VT \Rightarrow polymorphic VT and VF).
- * Mean age of onset 7-9yrs
- * Adrenaline provocation test / Exercise stress test
- * *RYR2* – autosomal dominant with high penetrance, 50-55%
- * *CASQ2* – autosomal recessive, 1-2%

CPVT

* Management

- * Avoid strenuous exercise / acute emotion
- * β -blockers
- * Implantable cardioverter defibrillator
- * FU by cardiologist
- * Screening of family members at risk

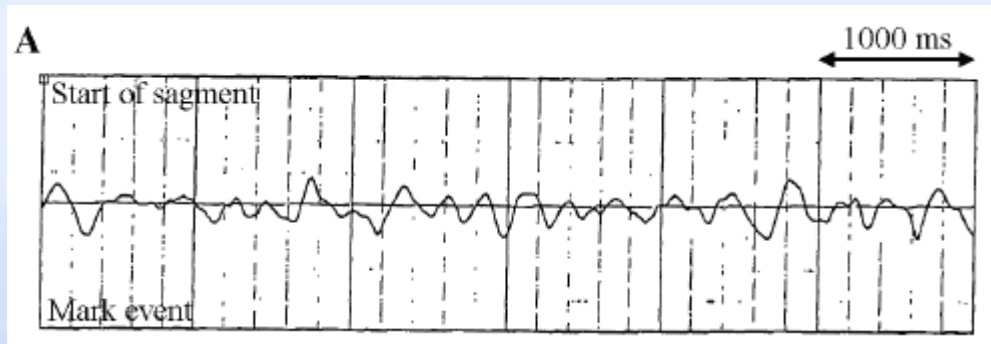


F/14

Good past health

Sudden collapse after boarding a bus. Regained consciousness shortly and collapsed again.

A similar episode 9 months earlier when quarrelling with a friend.



Baseline ECG – NAD

Echocardiogram – no structural heart disease

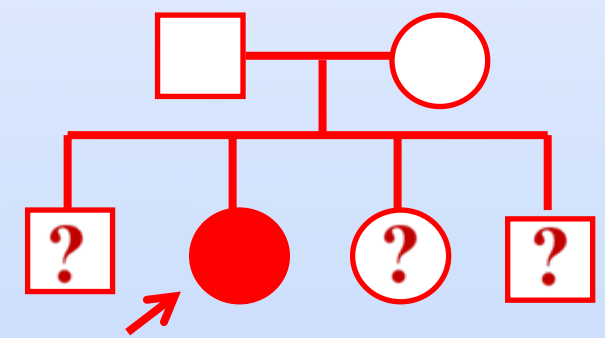
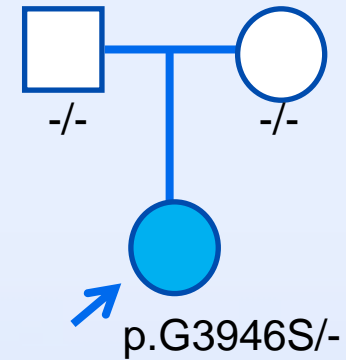
MR coronary angiogram – no anomalies of coronary arteries

CT brain – hypoxic brain damage

PVCs, polymorphic VT and non-sustained bidirectional VT when the patient was agitated.

Adrenaline provocation test

- * Family history –ve
- * *RYR2* mutation analysis
 - * 104 exons
 - * Critical regions
- * Heterozygous p.G3946S in exon 88
- * Both parents were negative
- * De novo mutation
- * Spastic paraplegia
- * Non-communicable
- * Bed-ridden



Molecular Autopsy

- * Postmortem genetic analysis
- * First report in 1999
- * *KCNQ1* mutation (LQT1) identified in a 19-yr-old woman who died after a near-drowning.

Definitive Dx for at least 60 relatives

Prophylactic therapy in mutation carriers

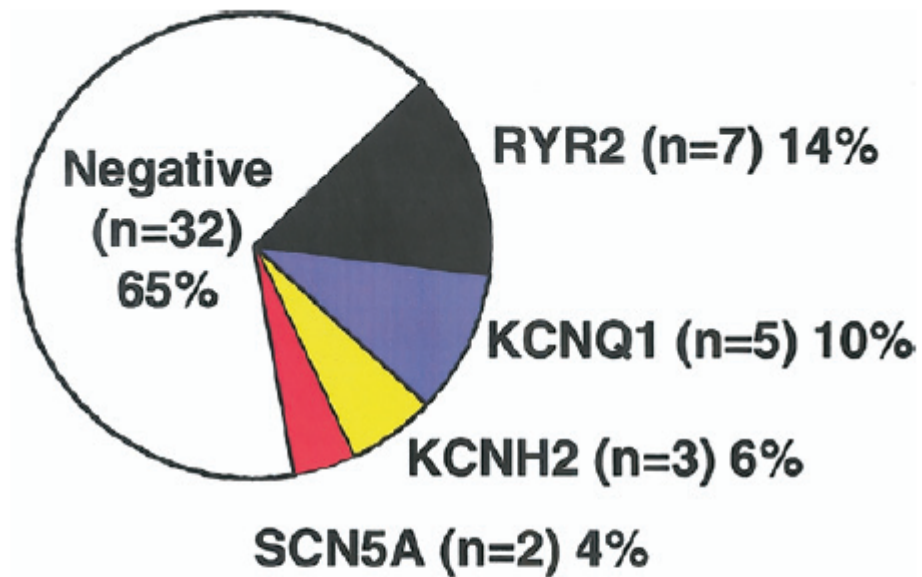


Figure 2

Summary of the Yield of Postmortem Cardiac Channel Genetic Testing in Cases of Autopsy-Negative SUD

Depicted is a pie chart summarizing the frequency and distribution of cardiac channel genotypes detected after a molecular autopsy of 49 medical examiner/coroner-referred cases of sudden unexplained death (SUD).

Cost-effectiveness analysis

- * Commercial LQTS genetic testing available in the US (US\$5400).
- * Symptomatic index cases who received a definite or inconclusive clinical diagnostic scores for LQTS.

	LQTS genetic testing	BRCA1/2 screening and oophorectomy	Mammography screening
Cost per year of life saved	\$2,500	\$3,900-\$1,600,000 for high- and average-risk women	\$21,400-\$117,680 for women ages 50-69yrs and 70-79yrs respectively.

-
- * *“More accurately diagnose and treat affected individuals.”*
 - * *“Clinicians can make more informed judgement about treatment.”*
 - * The cost and benefits of testing at-risk family members not included in the estimation \Rightarrow likely to be more favourable.

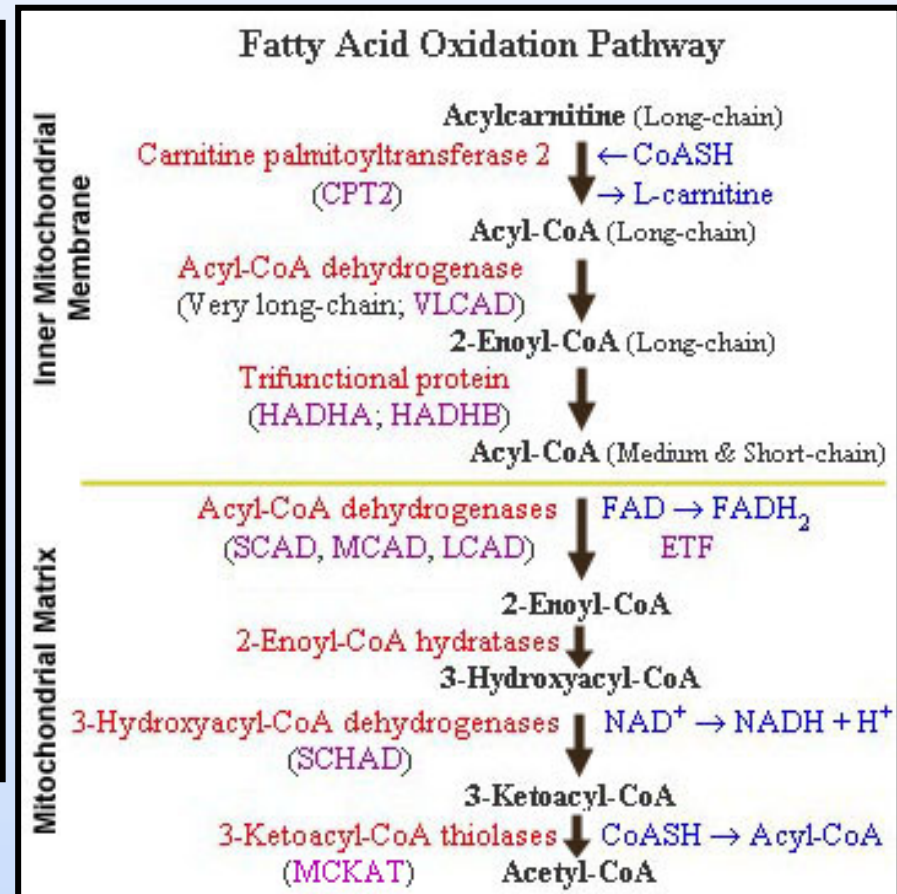
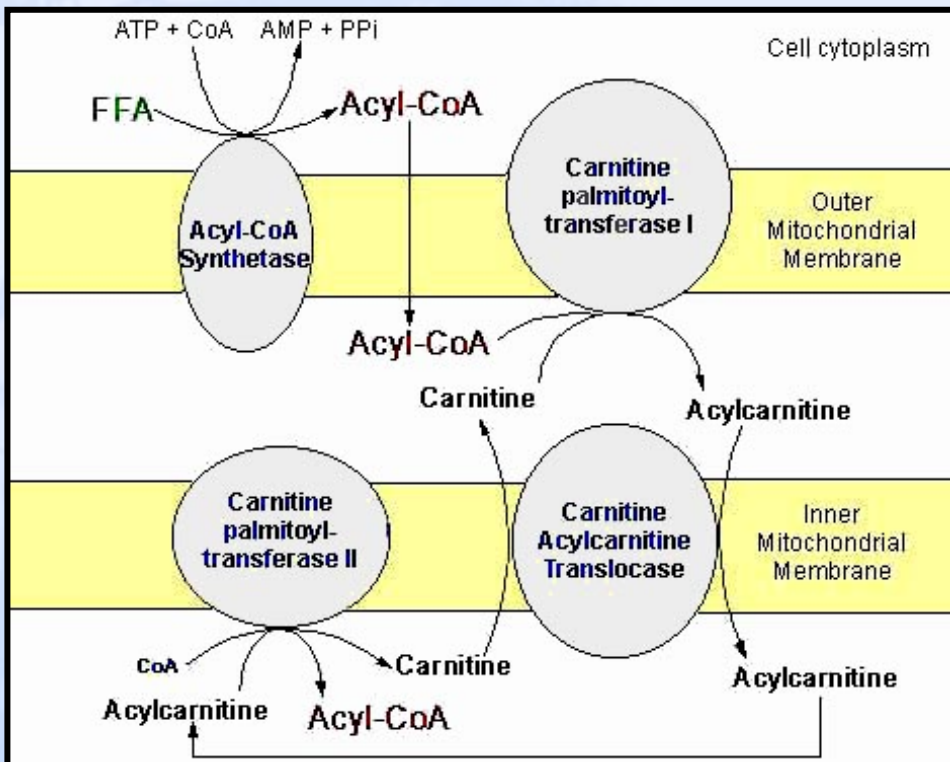
Challenges

- ✿ Genetic studies are expensive (time, manpower).
- ✿ Interpretation of results may not be straight-forward.
- ✿ How to prioritize patients?
- ✿ Genetic counselling



SUDDEN INFANT DEATH AND INBORN ERRORS OF METABOLISM

Fatty Acid Oxidation



CACT deficiency

- ✿ The 1st case in Hong Kong diagnosed in 2003
- ✿ M/3D, sudden death in hospital
- ✿ ~20 reported cases – sudden neonatal death is the commonest presentation.
- ✿ A lot of difficulties in the investigation process.

TABLE 1. Clinical features of the three patients diagnosed with carnitine-acylcarnitine translocase deficiency

Patient No.	Sex	Gestation (weeks)	Birth weight (kg)	Consanguinity	Feeding	Time at presentation (hours)	Initial symptoms	Survived
1	M	38.4	2.41	No	Formula	41	Sudden cardiac arrest	No
2	M	35.6	2.71	No	Breast-feeding	32	Sudden cardiac arrest	Yes
3	F	37.4	2.3	No	Formula	28	Cardiorespiratory failure	No

TABLE 2. Laboratory findings of patients with carnitine-acylcarnitine translocase deficiency at presentation and autopsy findings

Patient No.	Initial plasma glucose (mmol/L)	Maximum ammonia level (μ mol/L)	Long-chain acylcarnitine profiles	Dicarboxylic acid in urine	Lactate (mmol/L)	Autopsy
1	Not done	Not done	Not done	Raised	Not done	Steatosis in myocardium and hepatocytes
2	1.5	455	Raised	Raised	5.2	Not applicable
3	1.3	216	Raised	Raised	8.9	Not done

Founder mutation in CACT

- * *CACT* IVS2-10T>G
- * Homozygous in all 3 patients.
- * Reported previously in 2 Chinese parents of CACT patients (UK and USA).
- * Allow quick target mutation analysis.

Challenges

- ❖ Slow development in paediatric biochemical tests.
- ❖ Turnaround time cannot meet urgent need.
- ❖ Peri-mortem and post-mortem sampling protocol for IEM investigation.
- ❖ Genetic information alter the investigation protocol.
- ❖ Accumulation of expertise and sharing of clinical experience.

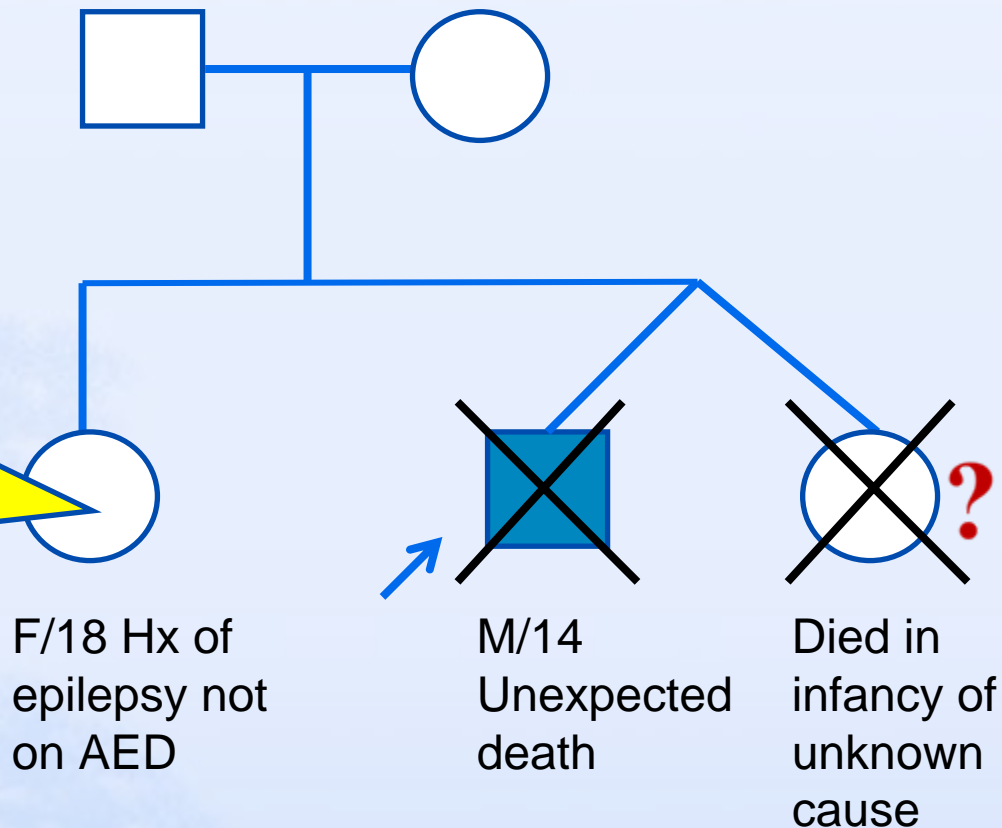


**UNEXPECTED DEATHS
CAUSED BY INBORN ERRORS
OF METABOLISM**

GAI

- * M/14
- * Presented to AED twice within 10hrs.
- * Acute deterioration after 9hrs of observation in AED.
- * Died 38hrs after admission.
- * No biological samples saved for metabolic investigation.
- * Retrospective family history – twin sister died of unknown cause in infancy.

**Abnormal
plasma
acylcarnitine!**



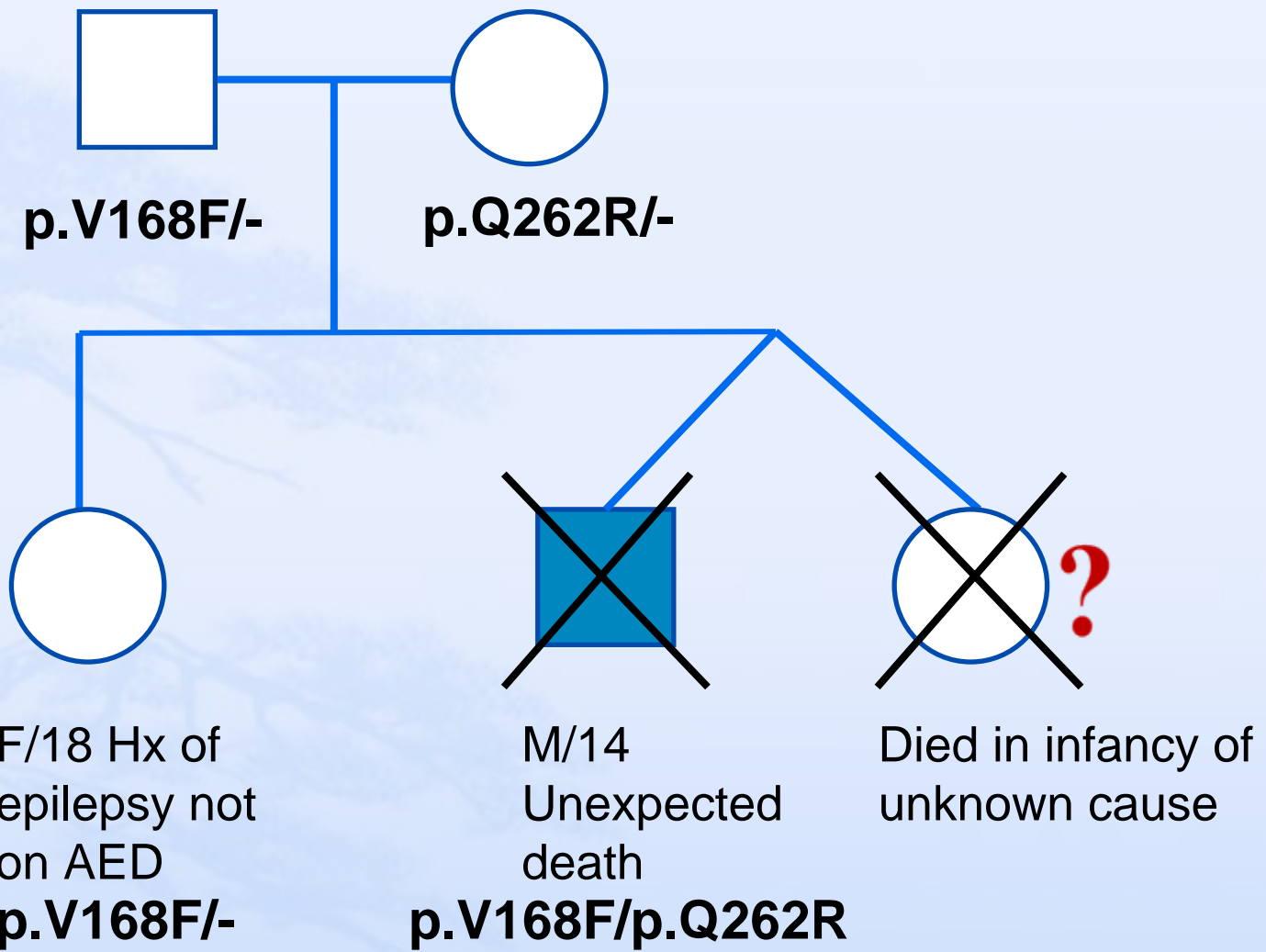
Urine: ↑ glutaric, 2-OHglutaric, ethylmalonic, isovalerylglycine, hexanoylglycine

Plasma: ↑ C4 to C16 acylcarnitines

⇒ Glutaric aciduria type II

- ✧ 3 known genes for GAI
 - ✧ *ETFA*
 - ✧ *ETFB*
 - ✧ *EFTDH*

- ✧ 2 Novel mutations in *ETFA*
 - ✧ p.V618F (paternal allele)
 - ✧ p.Q262R (maternal allele)



Challenges

- * GAI – a relatively common IEM in Hong Kong
- * Clinical suspicion
- * An inquest was held for this patient
 - * TAT of metabolic investigation.
 - * More comprehensive newborn screening program.

Conclusion

1. Increasing knowledge of genetic basis of **heritable** disorders which can cause sudden death / significant mortality and morbidity.
2. Increasing evidence which support of introduction of genetic testing into clinical practice.
3. Current situation in Hong Kong
 - a) No lack of clinical requests – lack of clinical guidelines
 - b) Piecemeal development – lack of central policy
 - c) Better support for up-stream and front-line investigations required.

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References

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