

ECG in AVB

- Predicting Torsades de Pointes

Jinhee Ahn, MD.

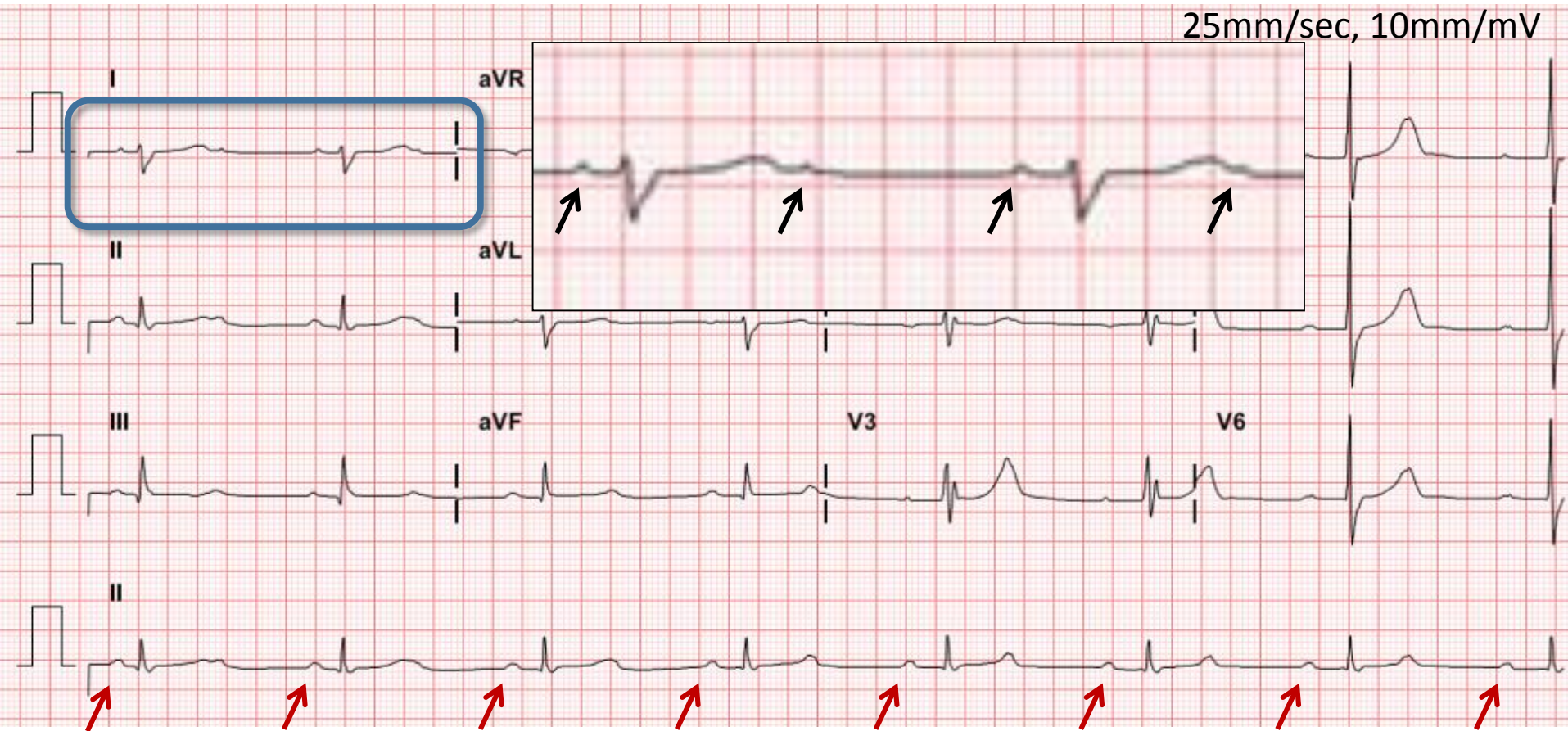
**Electrophysiology,
Division of Cardiology,
Pusan Nat'l Univ. Hosp.**



Case. Torsades de Pointes in CAVB

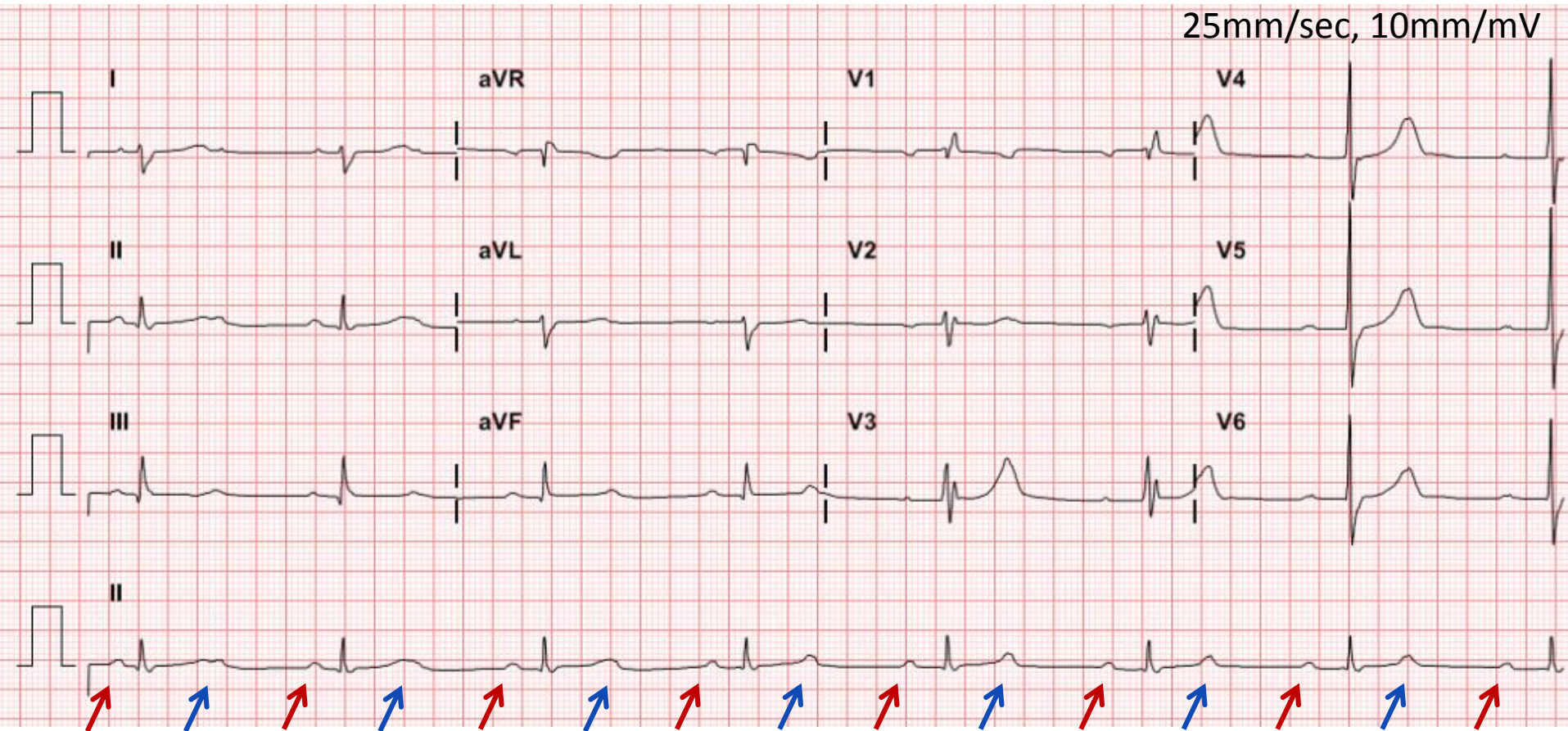
- F/78
- Underlying diseases
 - DM, HT, old CVA
- Spinal stenosis 수술 후 pneumonia 로 타원에서 supportive care 받던 분으로, dizziness 있어 본원 방문함.

내원시 심전도 (HR 44 bpm)



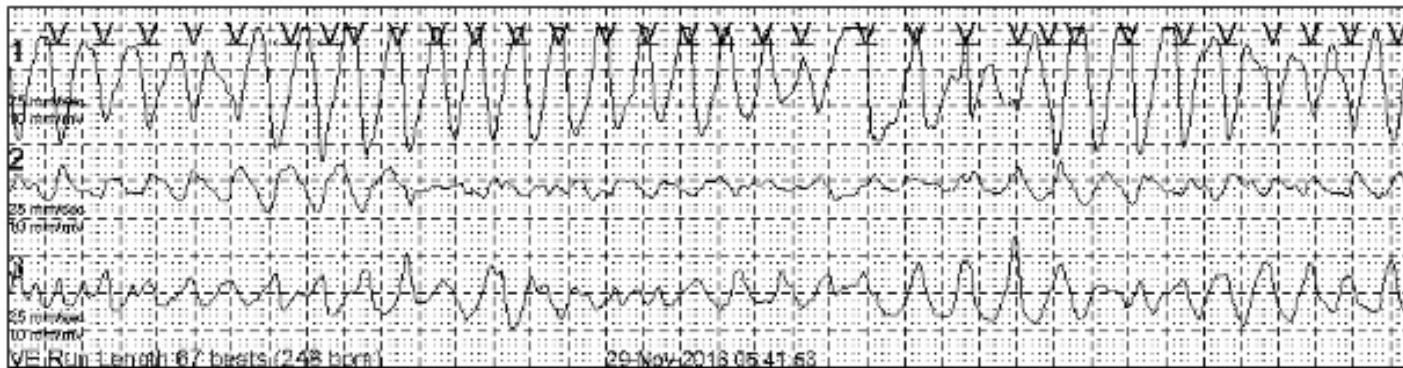
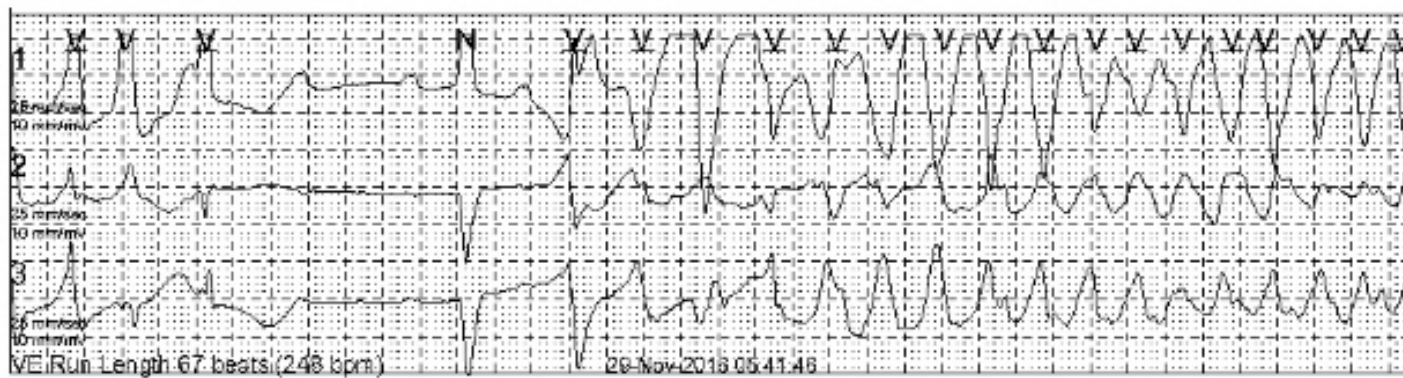
Sinus rhythm??

내원시 심전도 (HR 44 bpm)

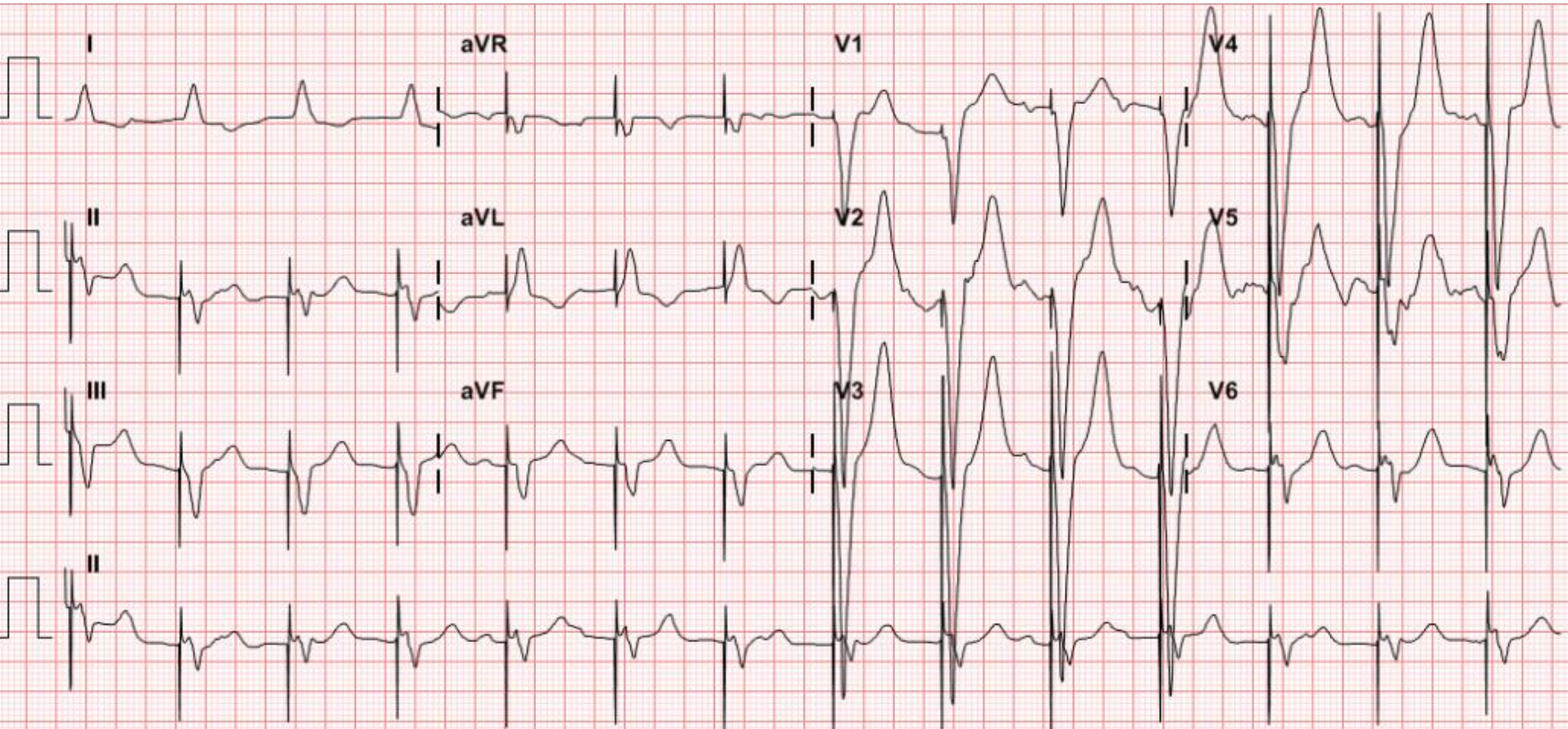


Complete AV block

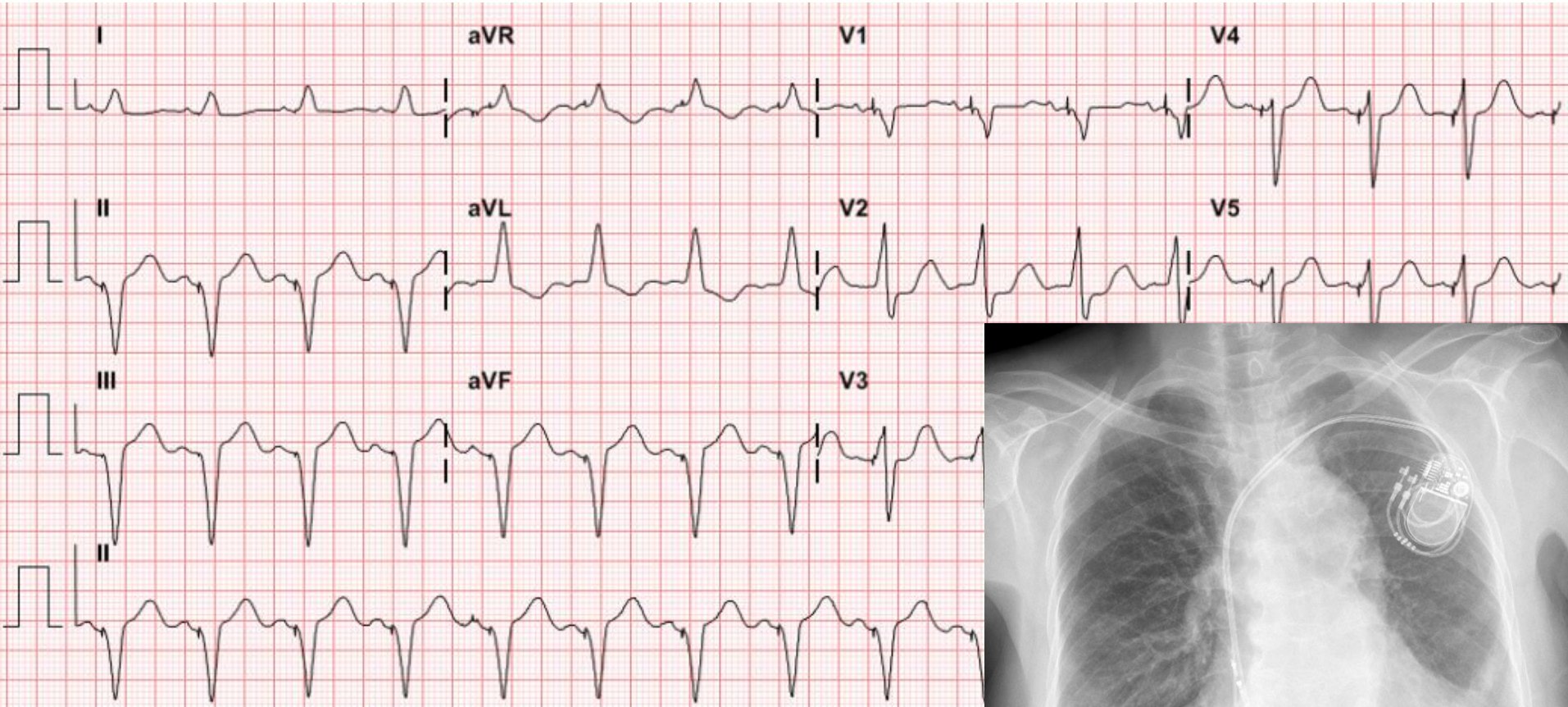
24hr Holter monitoring 중 syncope 발생



Temporary PM



Permanent PM implantation



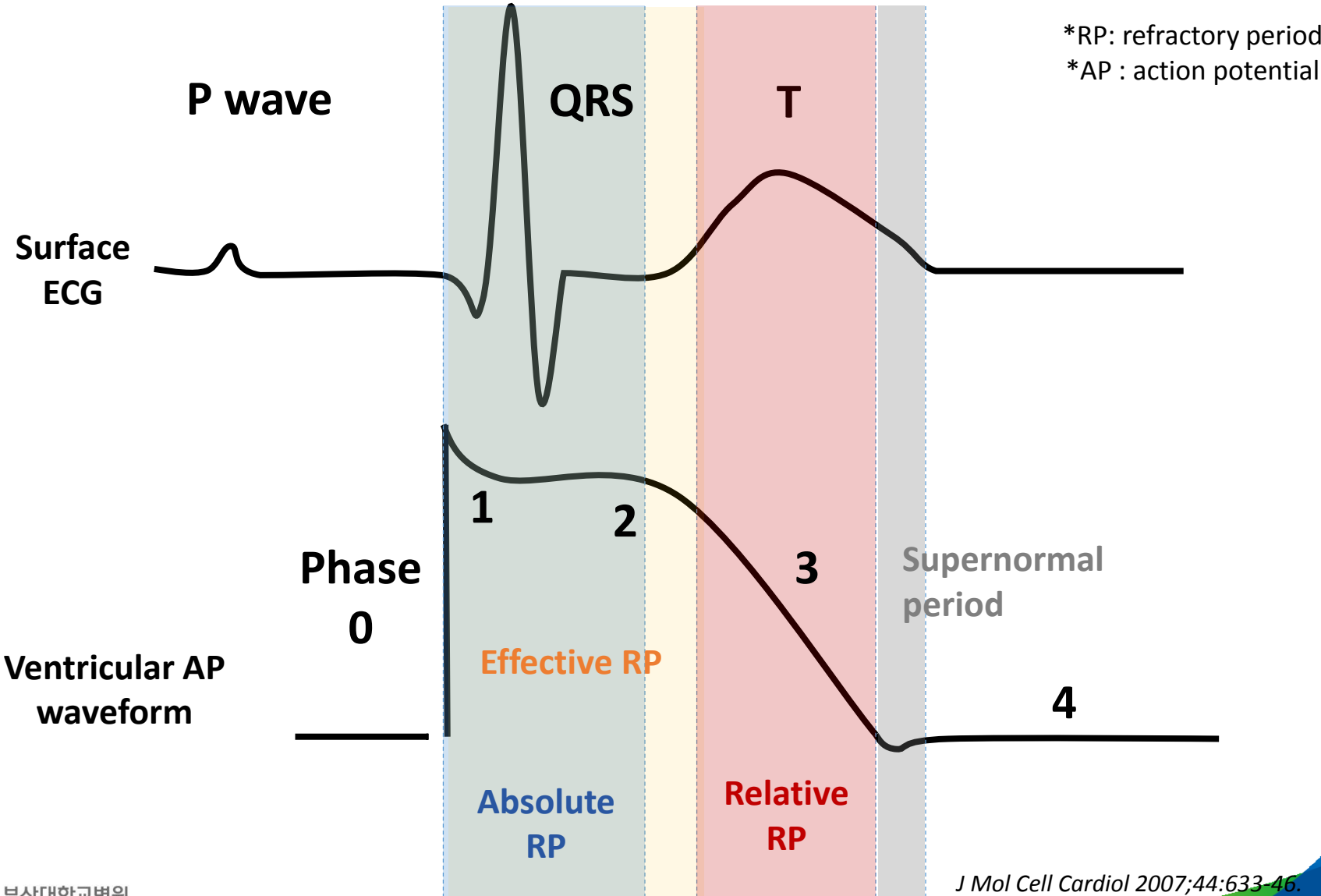
No more dizziness / syncope

What is Torsades de Pointes (TdP)?

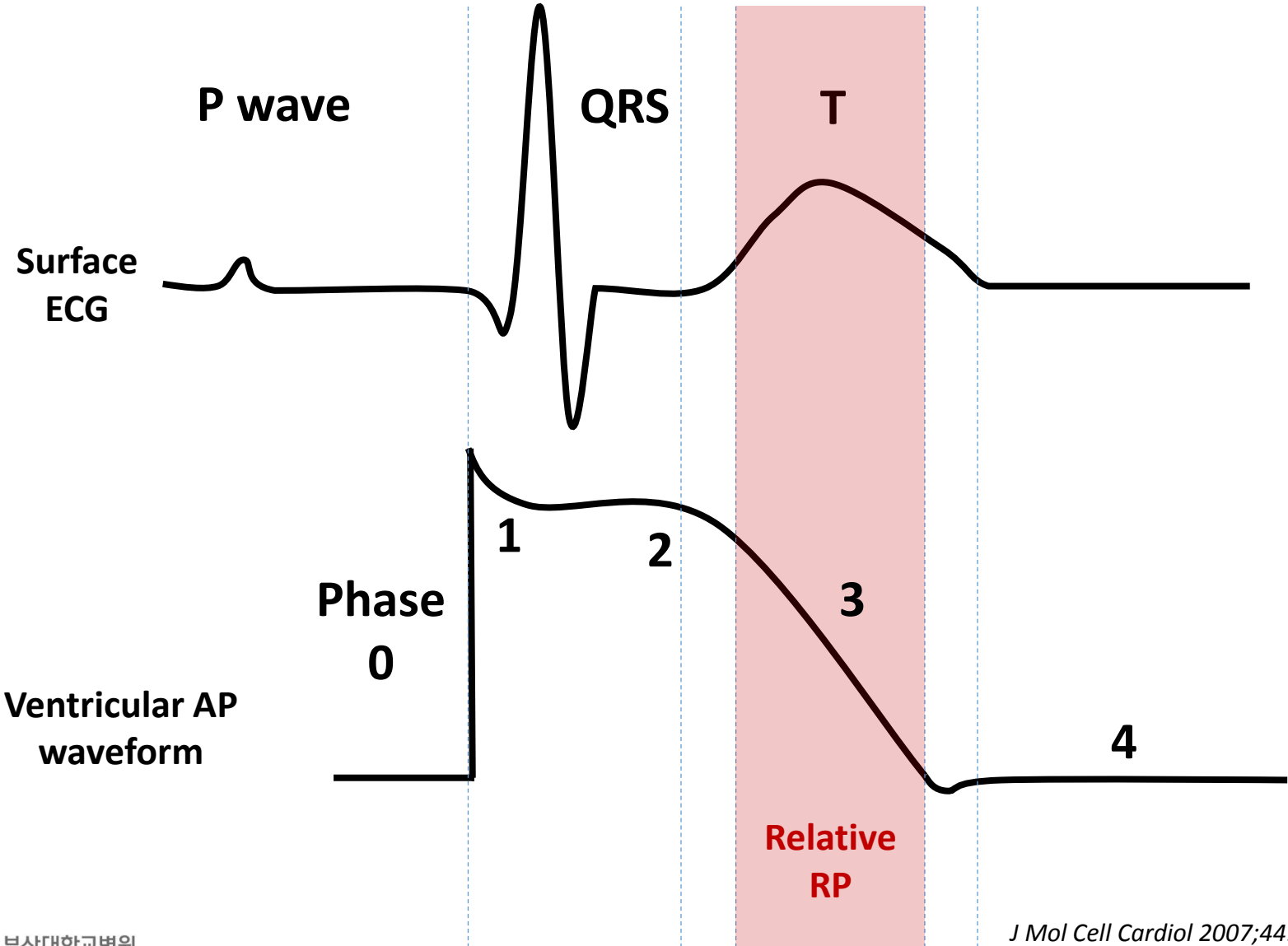
- A form of **PMVT** that occurs in the setting of **QT interval prolongation**.
- Terminologies to explain TdP
 - Long-short R-R cycle **sequences**
 - Transmural **dispersion** of repolarization
 - **QT prolongation** = prolongation of AP duration
 - **Early afterdepolarization**
 - PVC with **R on T** phenomenon
 - **Twisting** QRS complexes

Braunwald's heart disease. 10th ed. Vol. 1. pp782.

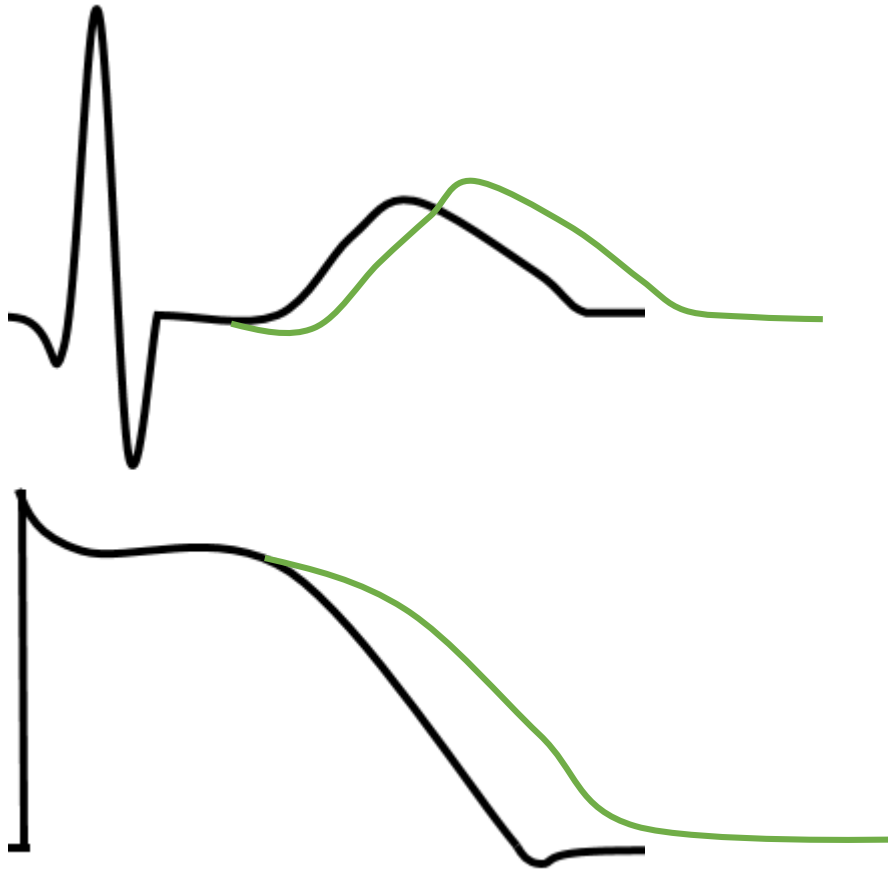
Relation btw Ventricular Action Potential and ECG



Relation btw Ventricular Action Potential and ECG



QT prolongation = prolonged AP duration



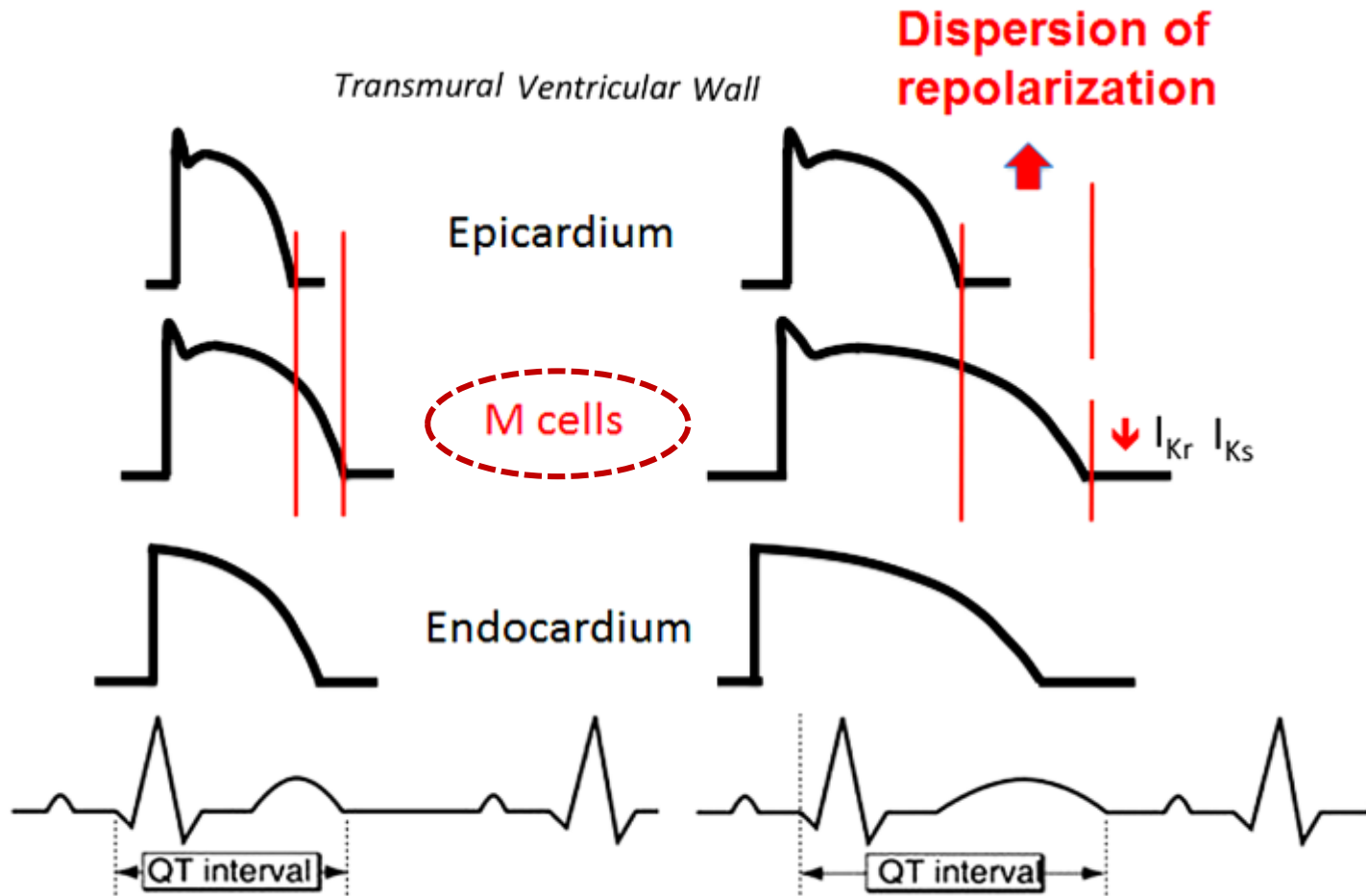
↓ Outward repolarizing
K current

or

↑ Inward depolarizing
Na or Ca current

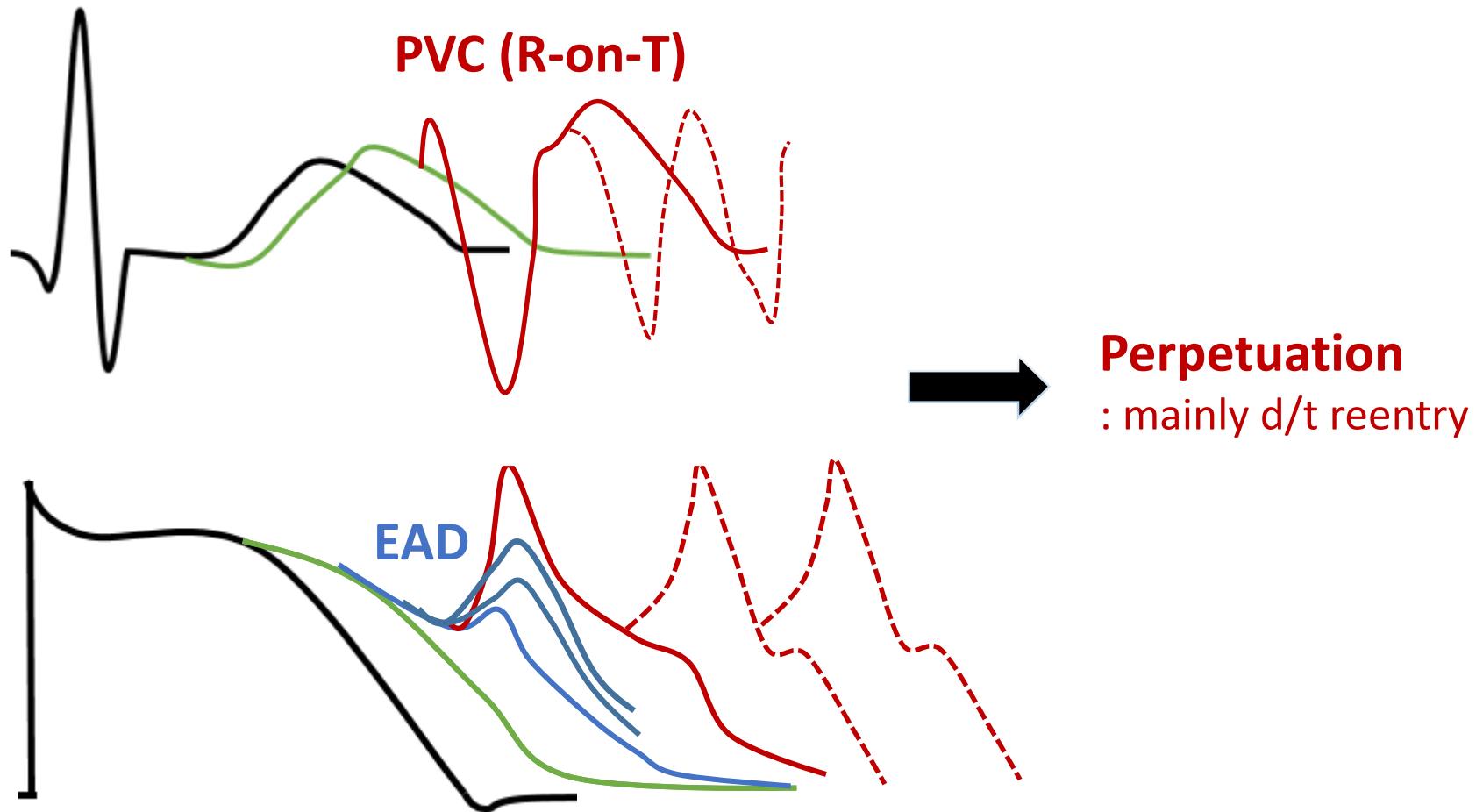
Transmural Dispersion of Repolarization

- AP duration difference between different myocardial layers
- Create a vulnerable window for development of reentry

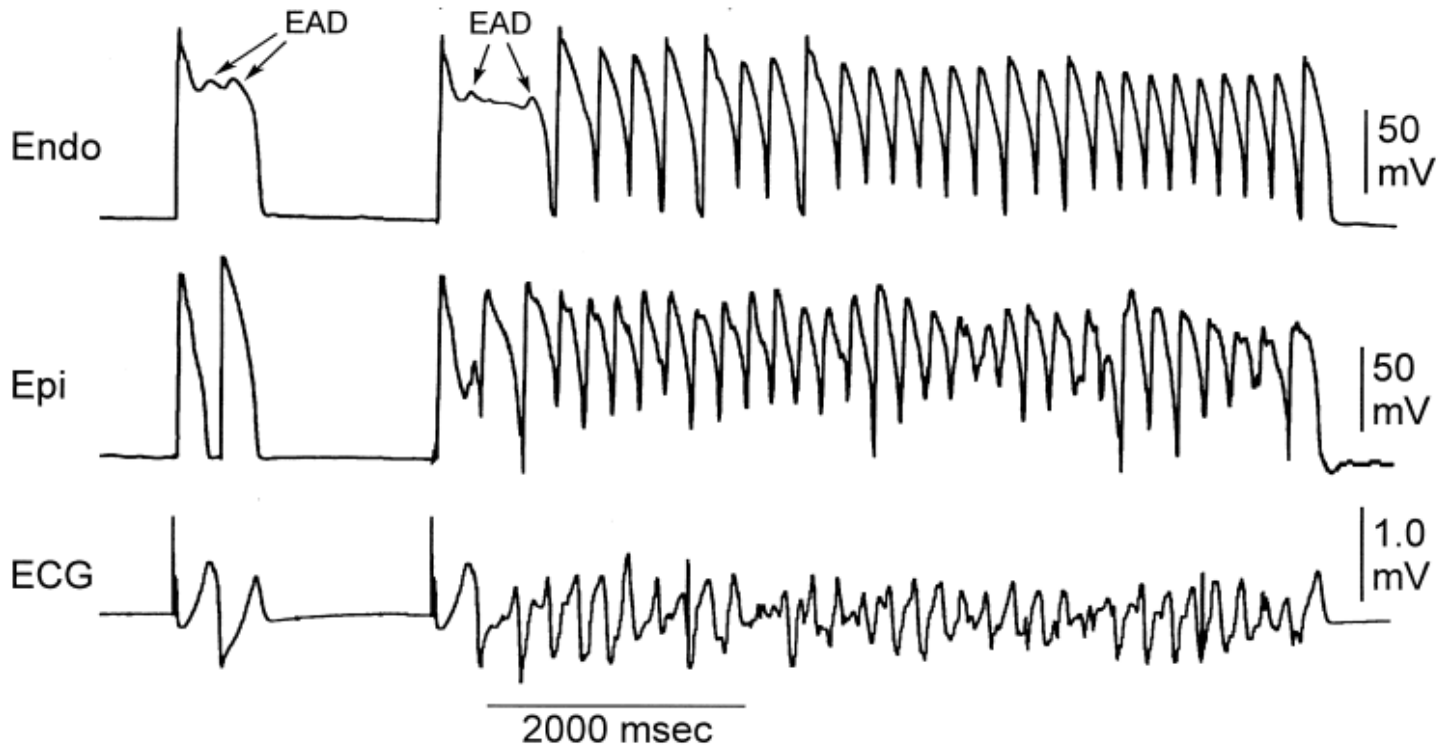


Early Afterdepolarization → PVC → TdP

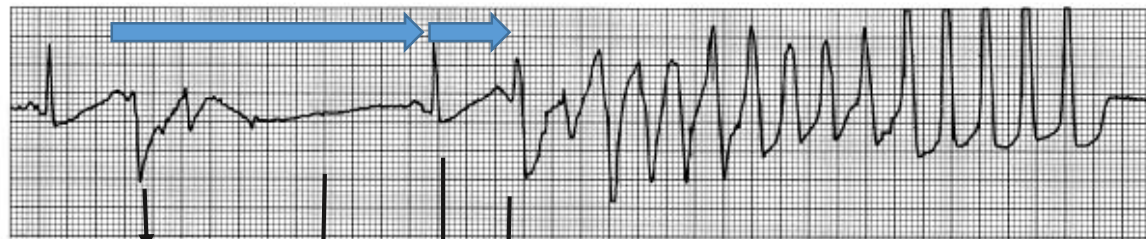
- Abnormal depolarization of Ca or Na → EAD → When reaching threshold potentials → generation of PVC (triggered beat) → initiation of TdP



TdP in an Isolated Arterially Perfused Rabbit Left Ventricle (dl-Sotalolol 100 μM)



TdP in a Patient with Prolonged QT Interval Who was on Sotalol



Long-short R-R sequence

PVC

Long
pause

EAD → PVC (R on T) triggering → TdP
Longer QT interval

Causes of TdP (QT prolongation)

1. Congenital long QT syndromes (see text for details)

Long QT syndrome type 1: Reduced repolarizing current I_{Ks} due to mutation in *KCNQ1* gene

Long QT syndrome type 2: Reduced repolarizing current I_{Kr} due to mutation in *KCNH2* gene

Long QT syndrome type 3: Delayed inactivation of the I_{Na} due to mutations in *SCN5A* gene

Others: Several other types of long QT syndromes have been described; long QT types 1, 2, and 3 account for 80–90% of cases

2. Acquired prolongation of QT interval

Electrolyte abnormalities

Hypokalemia

Hypomagnesemia

Hypocalcemia

Drugs

Antiarrhythmic drugs

Class IA: Quinidine, disopyramide, procainamide

Class III: Sotalol, amiodarone (QT prolongation common but torsade ventricular tachycardia is rare), ibutilide, dofetilide, almokalant

Antibiotics

Macrolides: Erythromycin, clarithromycin, azithromycin

Fluoroquinolones: Levofloxacin, moxifloxacin, gatifloxacin

Trimethoprim-sulfamethoxazole

Clindamycin

Pentamidine

Chloroquine

Antifungals: Ketoconazole, itraconazole

Antivirals: Amantadine

Antipsychotics

Haloperidol, phenothiazines, thioridazine, trifluoperazine, sertindole, zimelidine, ziprasidone

Tricyclic and tetracyclic antidepressants

Antihistamines (histamine 1-receptor antagonists)

Terfenadine, astemizole, diphenhydramine, hydroxyzine

Cholinergic antagonists: Cisapride, organophosphates

Citrate (massive blood transfusions)

Cocaine

Methadone

Fluoxetine (in conjunction with other drugs that prolong QT)

Cardiac conditions

Myocardial ischemia and infarction

Myocarditis

Marked bradycardia

Stress cardiomyopathy

Endocrine disorders

Hypothyroidism

Hyperparathyroidism

Pheochromocytoma

Hyperaldosteronism

Intracranial disorders

Subarachnoid hemorrhage

Thalamic hematoma

Cerebrovascular accident

Encephalitis

Head injury

Nutritional disorders

Anorexia nervosa

Starvation

Liquid protein diets

Gastroplasty and ileojejunal bypass

Celiac disease

ECG in AVB – factors predicting TdP

- Not everyone with AV block develops TdP.

Who are in a higher risk of TdP?

Clinical risk factors

- Older age
- Female gender
- HypoK, HypoCa, HypoMg
- Exposure to QT prolonging drugs
- Underlying disease
 - Heart failure
 - Left ventricular hypertrophy
 - Thyroid disease
 - Myocardial infarction
 - Obesity
- Polymorphisms or mutations in genes
- Various ECG findings

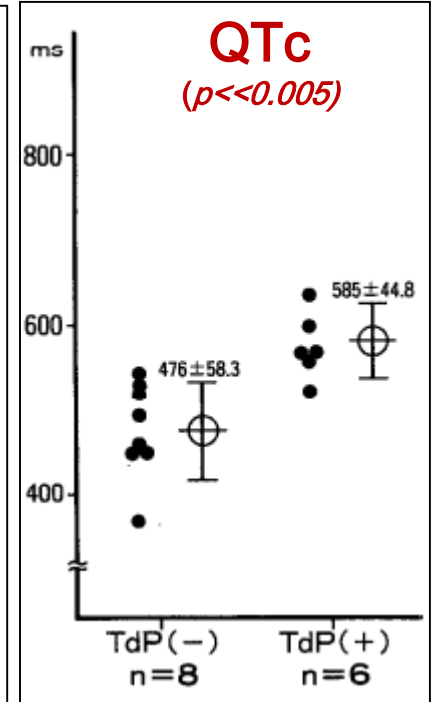
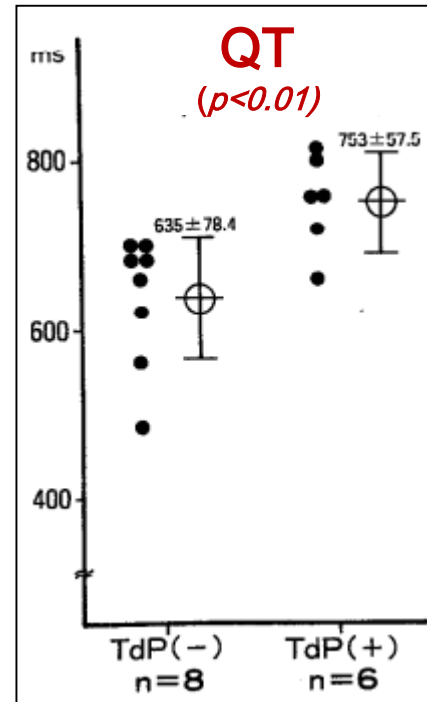
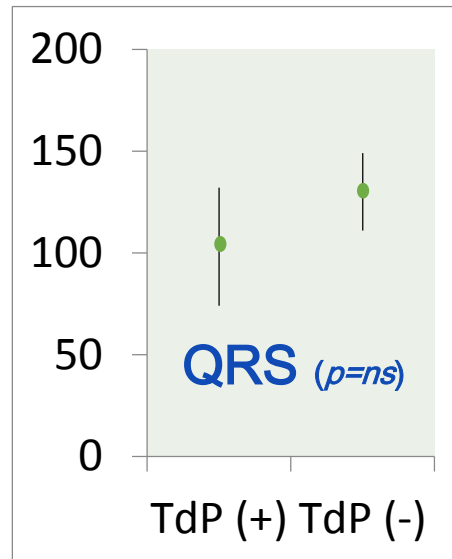
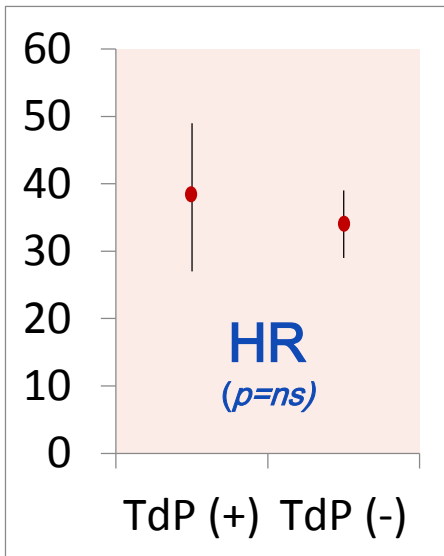
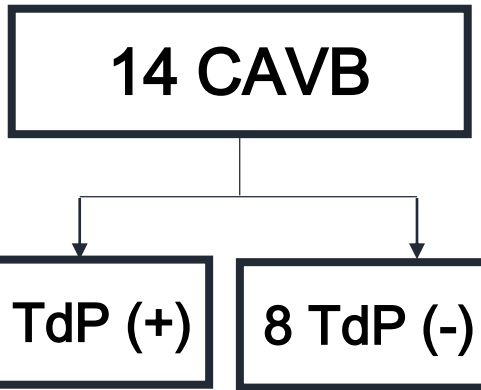
Eur Heart J 2003;24:1663-7.

J Intern Med 2006;259:59-69.

JAMA 1993;270:2590-7.

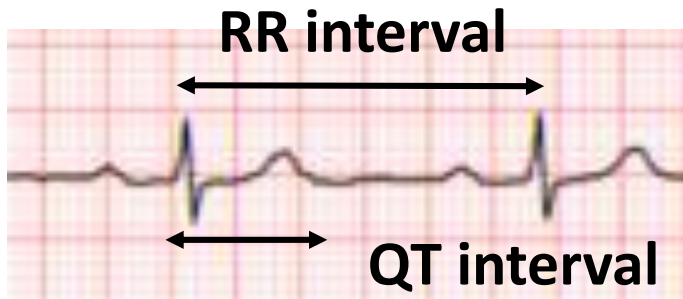
Eur J Cardiovasc Prev Rehabil 2005;12:363-8.

Bradycardia-Induced Abnormal QT Prolongation in Patients with Complete Atrioventricular Block with Torsades de Pointes



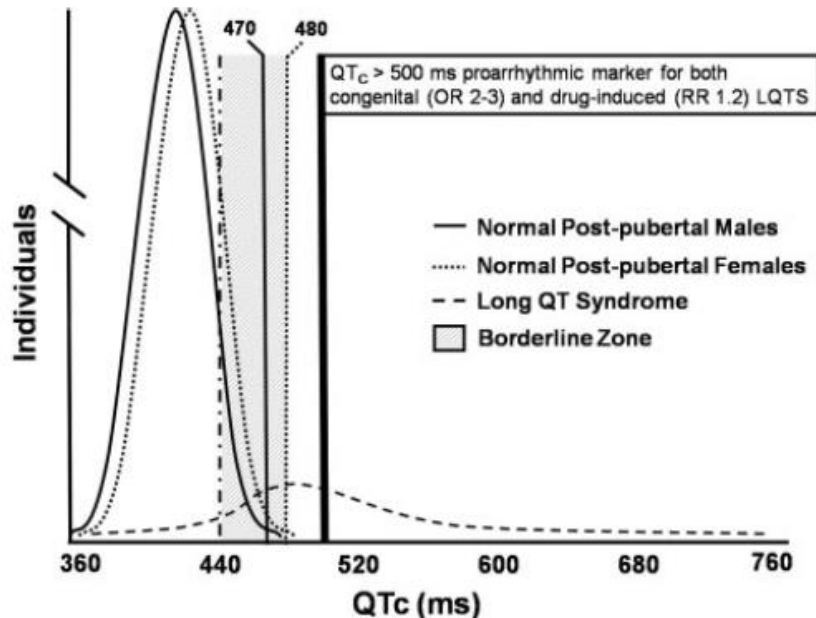
Am J Cardiol 1992;69:628-33.

a. QT (QTc) prolongation



$$QTc = QT \text{ interval} / \sqrt{RR}$$

Prevention of Torsade de Pointes in Hospital Settings A Scientific Statement From the American Heart Association and the American College of Cardiology Foundation



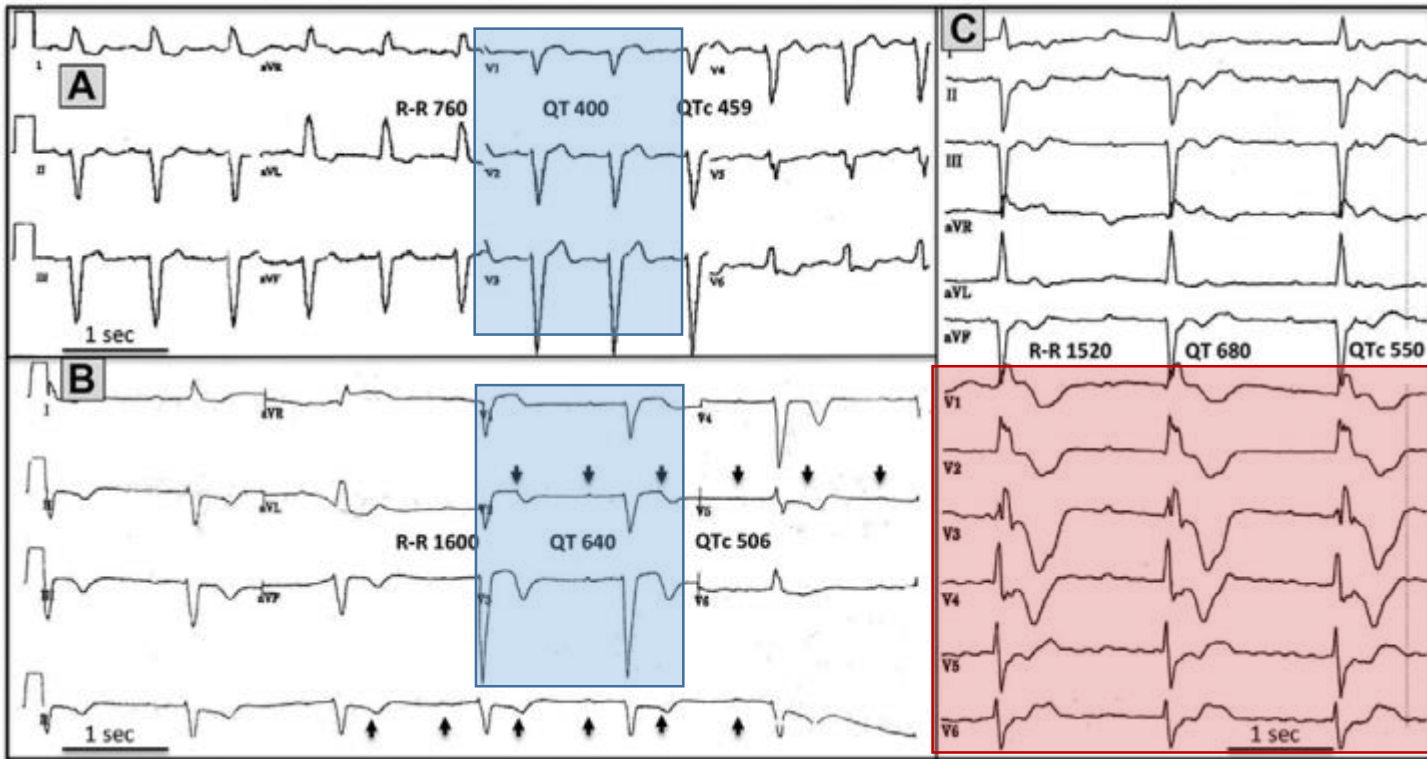
***Upper limits of normal (99th percentile) for QTc**
- 470ms (M), 480ms (F)

***QTc > 500ms**
- Proarrhythmic marker
Congenital (OR 2-3), acquired (OR 1.2)

Circulation 2010;121:1047-60.

b. QRS morphology change

Long QT Syndrome Complicating Atrioventricular Block Arrhythmogenic Effects of Cardiac Memory

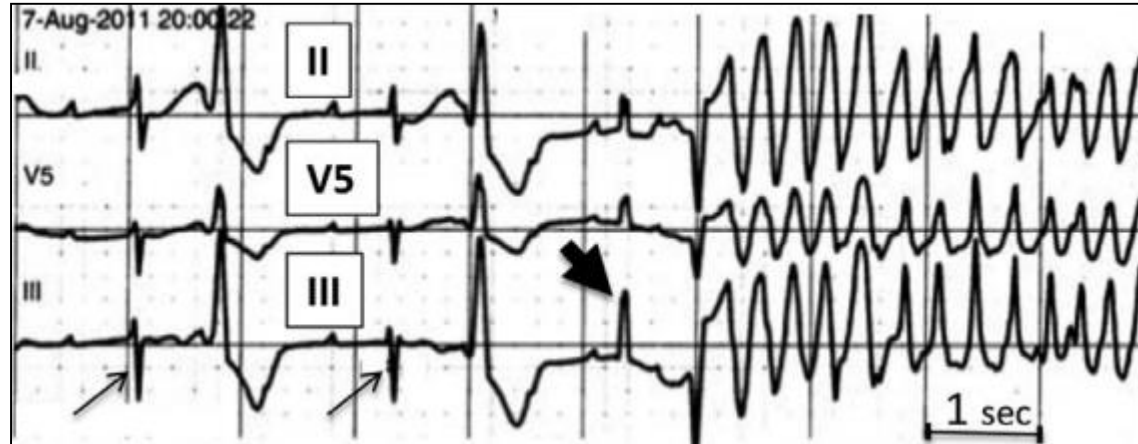


CAVB
(VR 38bpm)

2:1 AVB
(VR 36bpm)

Long QT Syndrome Complicating Atrioventricular Block

Arrhythmogenic Effects of Cardiac Memory



Circ AE 2014;7:1129-35.

Holter of our case patient



Long QT Syndrome Complicating Atrioventricular Block

Arrhythmogenic Effects of Cardiac Memory

- A change in QRS morphology at the time of AVB – in **1/3 patients**
- T wave change, QT prolongation d/t **cardiac memory** → TdP risk

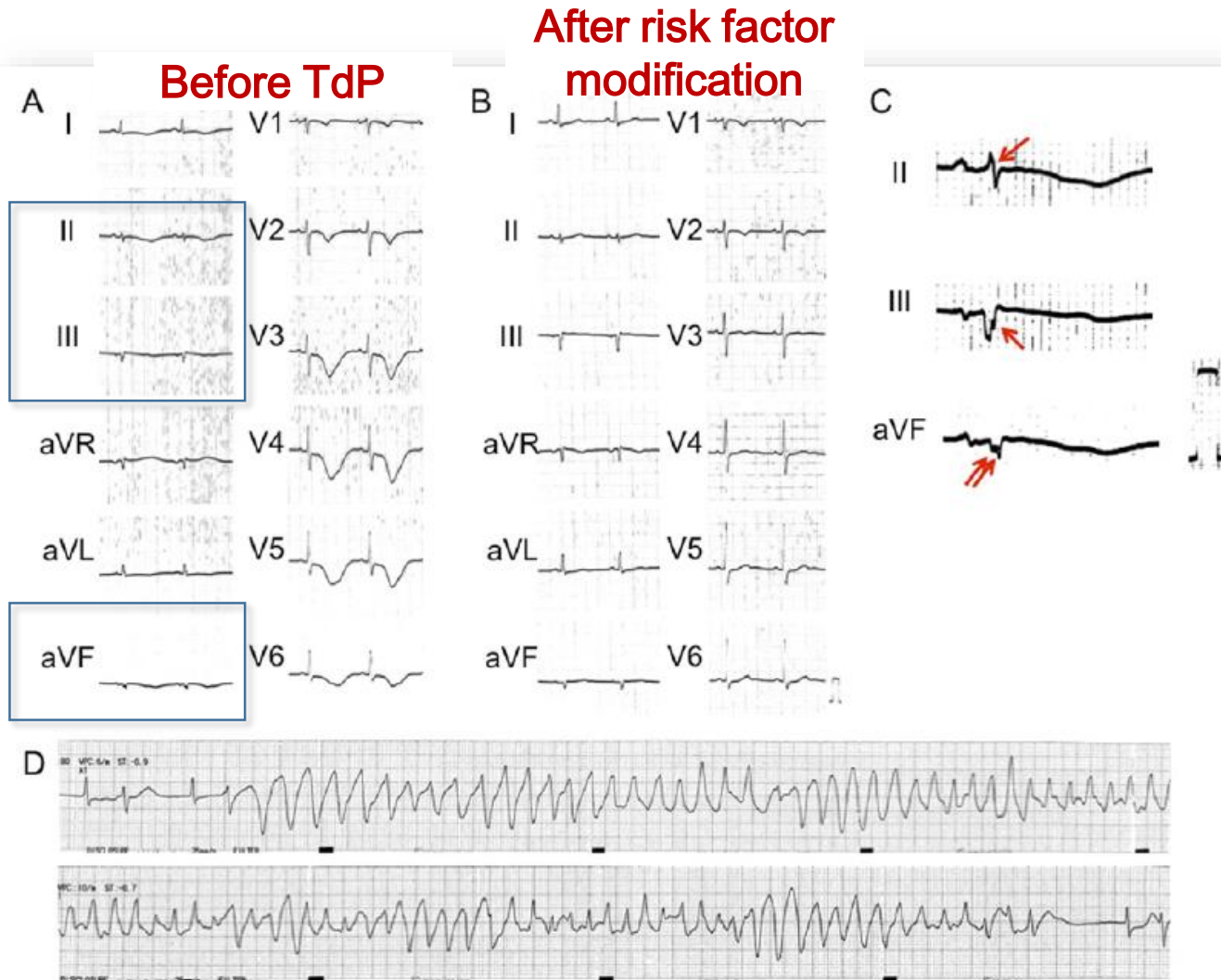
| | No Change in QRS Axis | QRS Axis Change | P Value |
|--------------------|-----------------------|-----------------|---------|
| ECG during AVB† | | | |
| QTc | 442±62 | 501±89 | 0.001 |
| QTcF | 479±63 | 543±88 | <0.001 |
| ΔQT | 153±82 | 222±96 | <0.001 |
| ΔQTc | -5±68 | 43±91 | 0.006 |
| Long QT‡ | | | |
| QT≥570 ms | 31 (45%) | 27 (73%) | 0.008 |
| QTc≥480 ms | 19 (28%) | 22 (60%) | 0.002 |
| Torsade de pointes | 4 (6%) | 8 (22%) | 0.023 |

| Predictor of TdP | Odds Ratio (95% Confidence Intervals) | P Value |
|---|---------------------------------------|--------------|
| Model 1 (without QTc) | | |
| Female sex | 9.6 (1.5–59.9) | 0.016 |
| Complete AVB* | 2.1 (0.1–23.1) | 0.54 |
| R-R during AVB (for 100-ms increase) | 1.1 (0.9–1.3) | 0.46 |
| QRS width during AVB (for 10-ms increase) | 0.9 (0.7–1.0) | 0.06 |
| Change in QRS morphology | 5.4 (1.4–21.3) | 0.016 |
| Change in QRS axis | 3.1 (0.7–13.5) | 0.130 |
| Change in QRS morphology and axis | 10.7 (2.0–57.6) | 0.006 |

Circ AE 2014;7:1129-35.

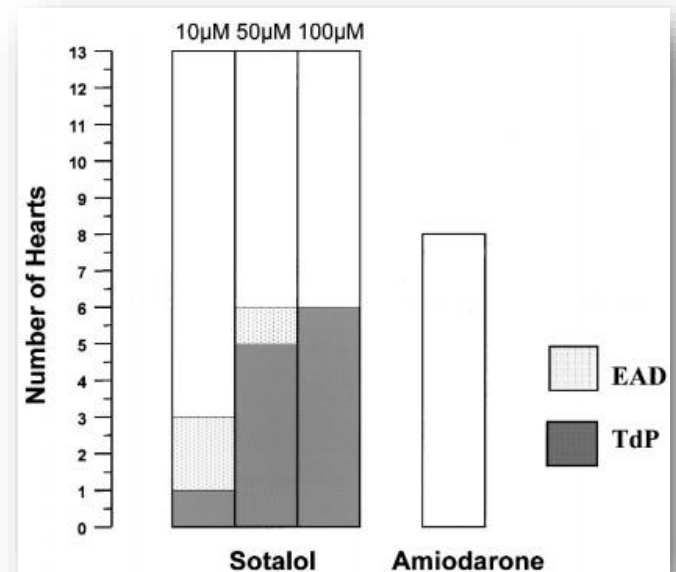
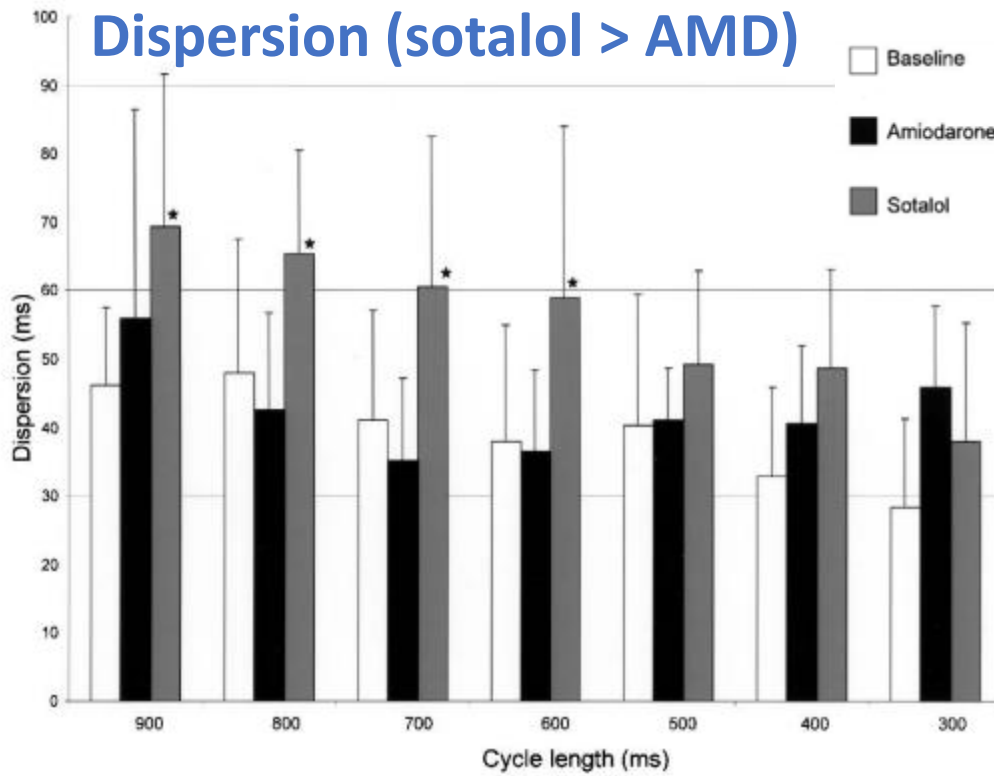
Fragmented QRS is associated with torsades de pointes in patients with acquired long QT syndrome

- **Fragmented QRS** - Depolarization abnormality
 - Substrate for reentrant circuit after TdP initiation



c. Transmural dispersion of repolarization

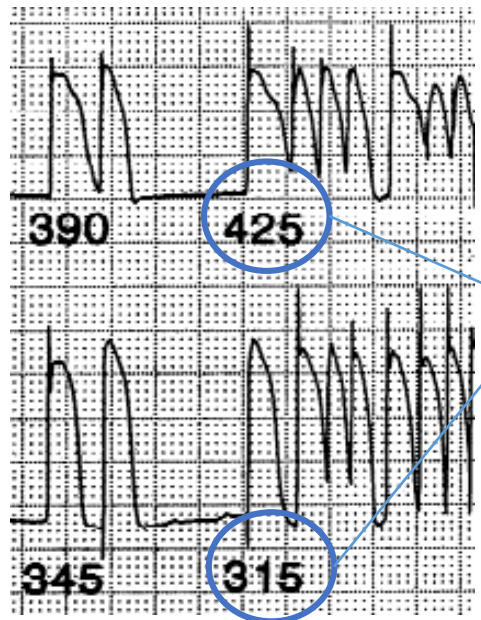
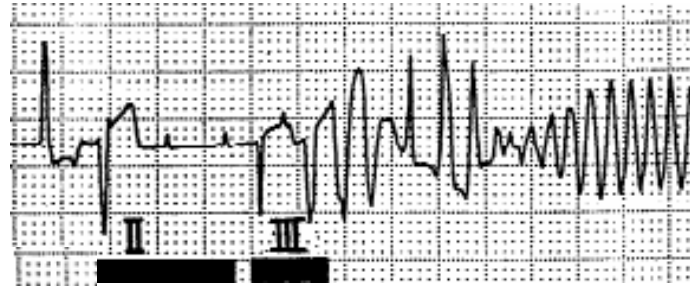
- An increase in QT interval does not necessarily lead to TdP.
- Comparison of TdP induction between class III AAD.
 - In a rabbit model of acute AV block
 - **sotalol vs. amiodarone**



EAD, TdP (sotalol > AMD)

c-1. Interventricular dispersion of repolarization

- Long-short sequence



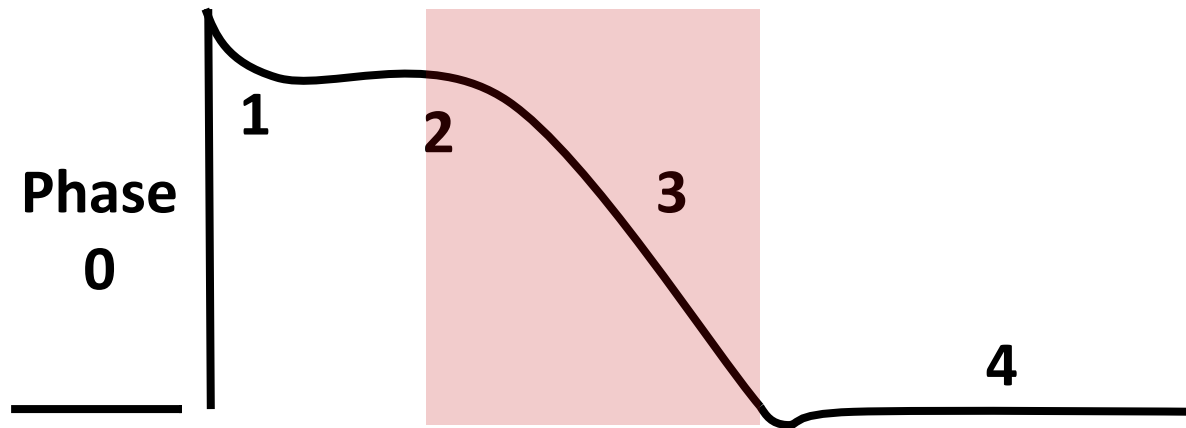
LV APD

RV APD

Interventricular dispersion of repolarization (110ms) → TdP

d. T wave morphology

- Prolongation of AP duration
- Marked transmural dispersion of repolarization
- EAD triggering PVC, perpetuating TdP



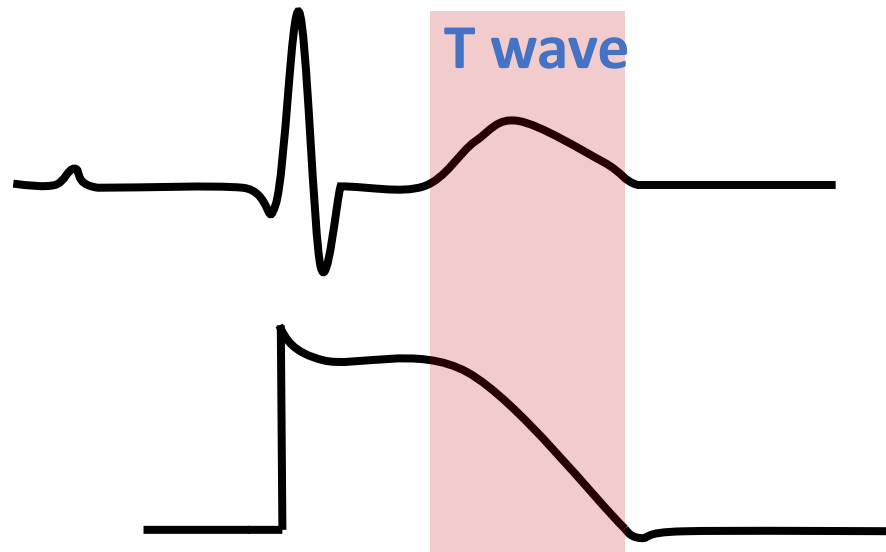
Repolarization problem



QT interval prolongation

T wave morphologic change

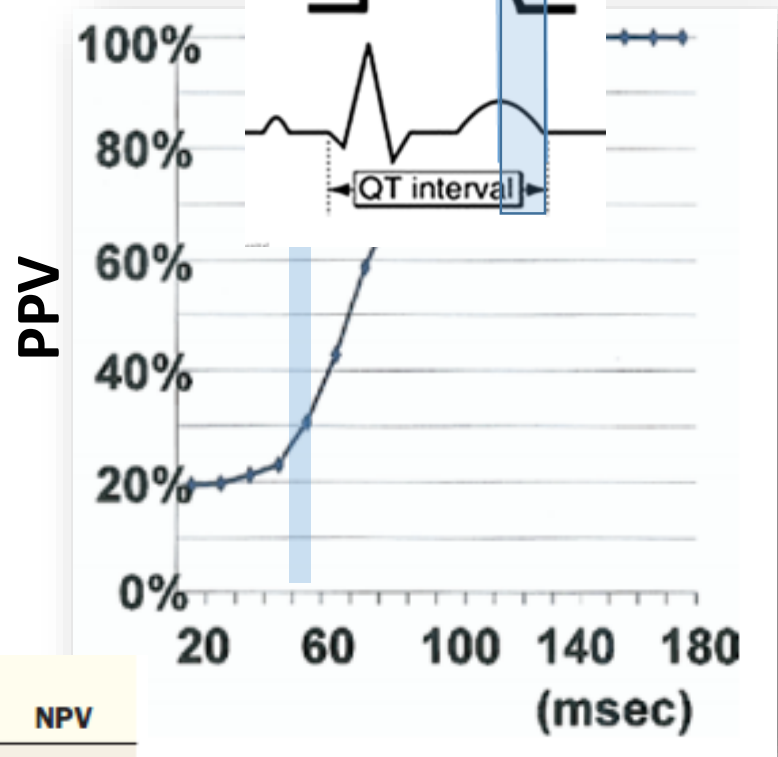
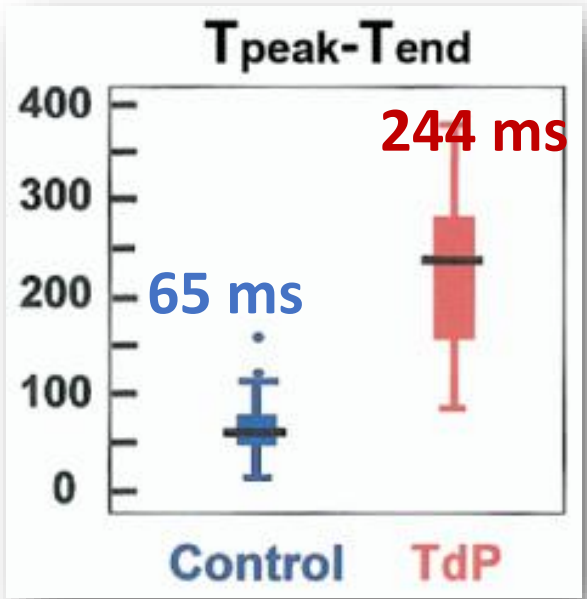
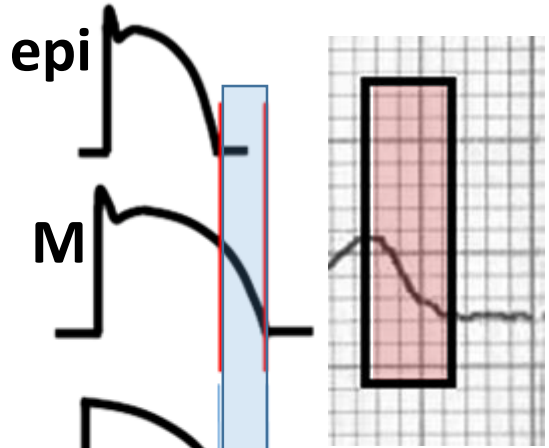
d. T wave morphology



- T [peak] – T [end] interval
- Notched T wave (similar to LQT2)
- Complex (triphasic) T wave
- T wave alternans (macroscopic beat-to-beat alteration)
- Long QT with abnormal T-U wave

d-1. T [peak] – T [end] interval

- Comparison of Tpeak –Tend
- Pts with CAVB (TdP (-) vs (+))



| Parameter | Cutoff Value | Sensitivity | Specificity | PPV | NPV |
|-------------------------------------|--------------|-------------|-------------|-------|-------|
| T _{peak} -T _{end} | 117 ms | 96.6% | 98.2% | 93.3% | 99.1% |

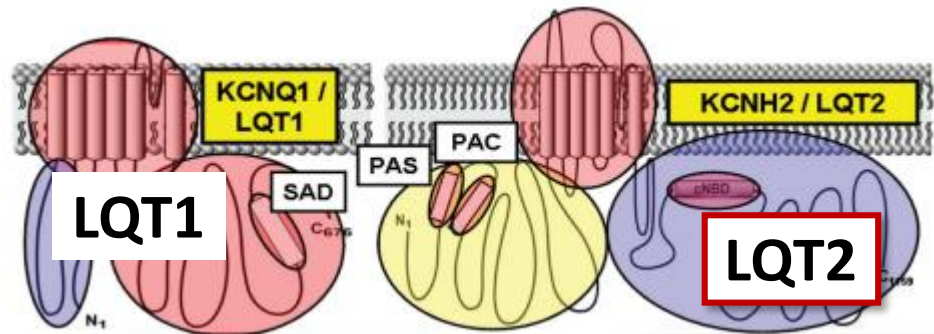
d-2. Notched T wave (similar to LQT2)

Cardiomyocyte Action Potential

Phase 3 : blocking delayed rectifier K current
 → QT prolongation

I_{ks} (KCNQ1)

I_{kr} (KCNH2)



Broad T



Bifid T

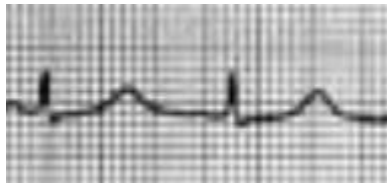
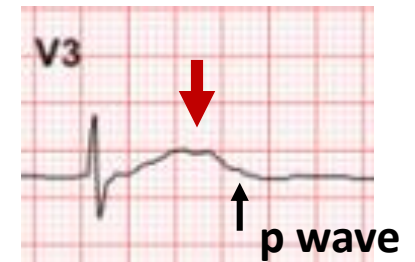
| | |
|---|---|
| 0 | Na ⁺ (in) ✓ I _{Na} (rapid) |
| 1 | K ⁺ , Cl ⁻ (out) ✓ I _{to1,2a} (transient outward) |
| 2 | Ca ²⁺ (in), K ⁺ (out) ✓ I _{Ca-t} (Ca long) ✓ I _{KS} (K slow delayed rect.) |
| 3 | K ⁺ (out) ✓ I _{KS} (K slow delayed rect.) ✓ I _{KR} (K rapid delayed rect.) ✓ I _{K1} (inward rect.) |
| 4 | K ⁺ (out) ✓ I _{K1} (inward rect.) |

LQT2-like T wave morphology

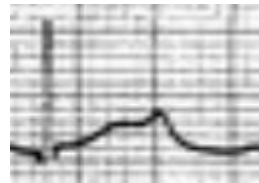
Table 1

Clinical and Electrocardiographic Characteristics of Patients With Bradyarrhythmias With and Without Torsade de Pointes

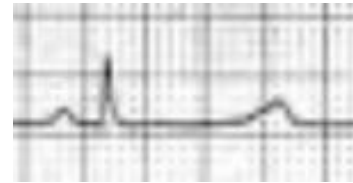
| Variable | TdP n = 30 | Control n = 113 | p Value |
|----------------------|-------------------|--------------------|------------------|
| QT morphology | | | |
| LQT1-like | 4 (13.8%) | 10 (8.8%) | 0.426 |
| LQT2-like | 16 (55.2%) | 3 (2.7%) | <0.001 |
| LQT2 with T2>>T1 | 8 (27.6%) | 0 (0%) | <0.001 |
| LQT3-like | 2 (6.9%) | 6 (5.3%) | 0.741 |
| "Bumps-ahead sign" | 17 (58.6%) | 20 (17.7%) | <0.001 |



LQT1



LQT2



LQT3

Torsades de pointes complicating atrioventricular block: Evidence for a genetic predisposition

Tested gene

: HERG, KCNQ1, KCNE1, KCNE2,
SCN5A

29 with CAVB, QT>600ms

vs.

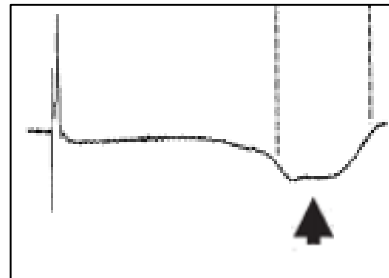
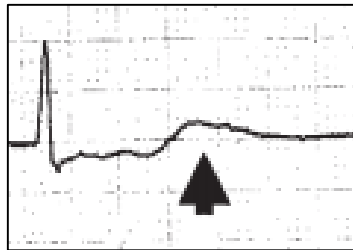
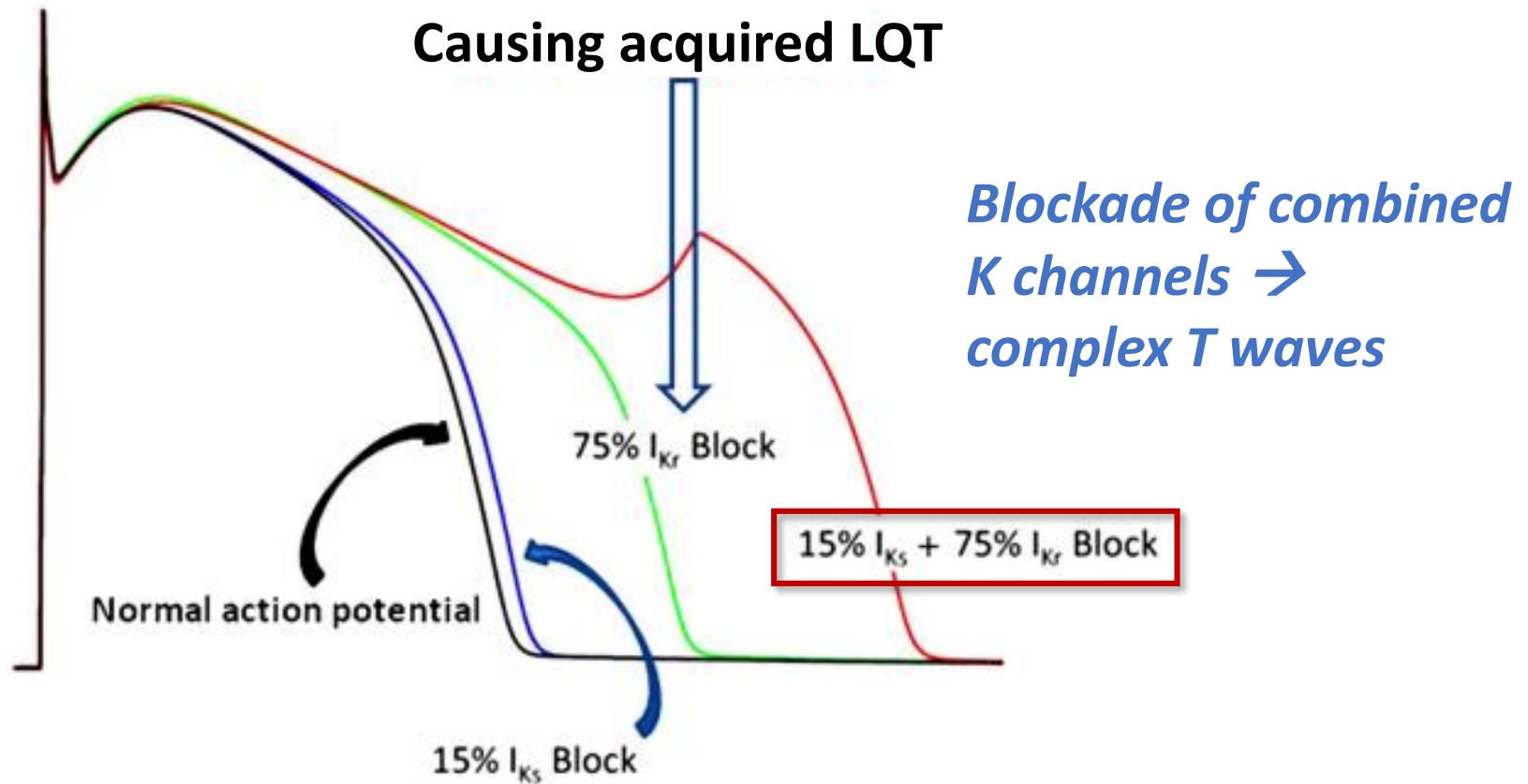
22 with CAVB, QT<600ms

vs.

100 healthy controls

CAVB with LQT was
associated with **HERG**
mutation in 17% !

d-3. Complex T wave



d-3. Complex (Triphasic) T wave

Table 1

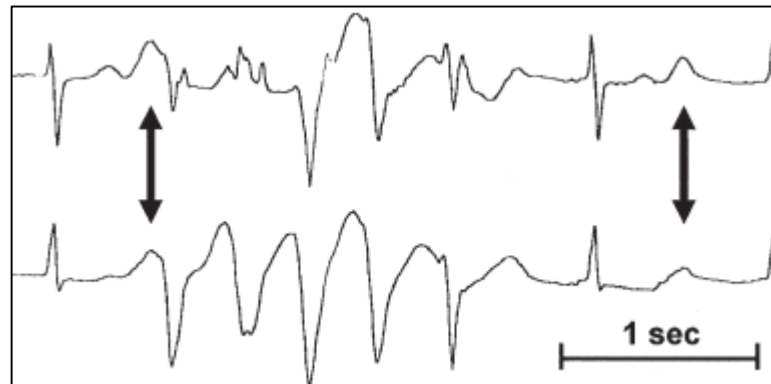
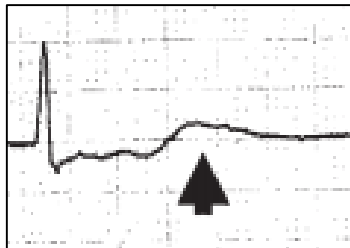
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| "Bumps-ahead sign" | 17 (58.6%) | 20 (17.7%) | <0.001 |



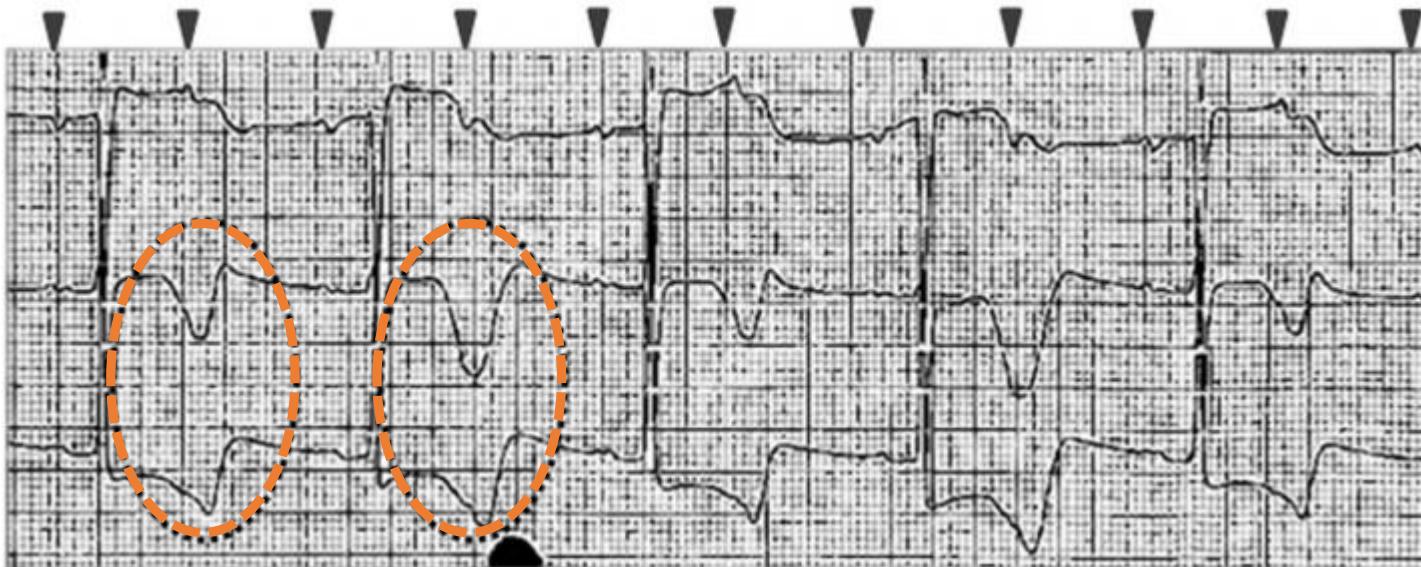
U wave?
Sign of EAD?

Triphasic T : late positive component

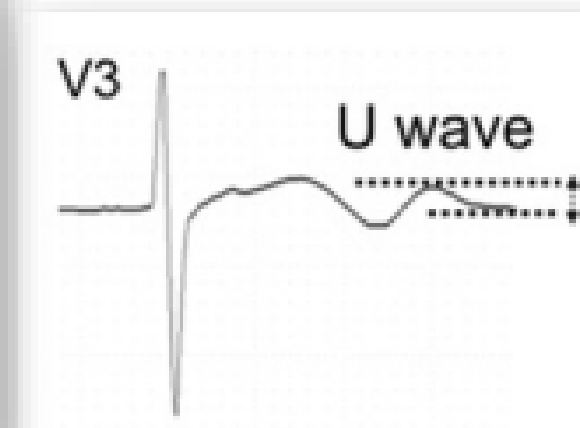
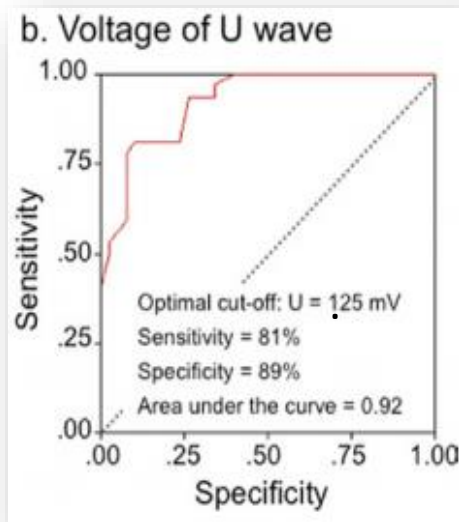


d-4. Others

- T wave alternans (macroscopic beat-to-beat alteration)



- Long QT wave with abnormal T-U wave



A 2-Step Model for predicting TdP

The Morphology of the QT Interval Predicts Torsade de Pointes During Acquired Bradyarrhythmias

| Parameter | Cutoff Value | Sensitivity | Specificity | PPV | NPV |
|----------------------------------|--------------|-------------|-------------|-------|-------|
| $T_{\text{peak}}-T_{\text{end}}$ | 117 ms | 96.6% | 98.2% | 93.3% | 99.1% |
| QTc interval | 480 ms | 96.6% | 92.0% | 75.7% | 99.0% |
| QT interval | 570 ms | 90.0% | 86.7% | 64.3% | 97.0% |

Values derived from receiver-operating characteristic curves.

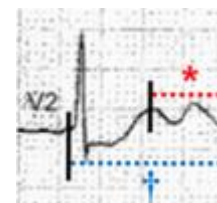
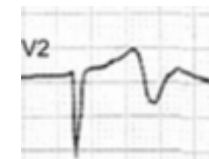
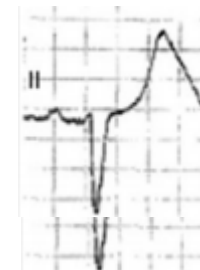
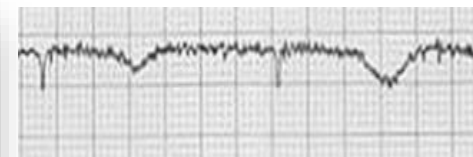
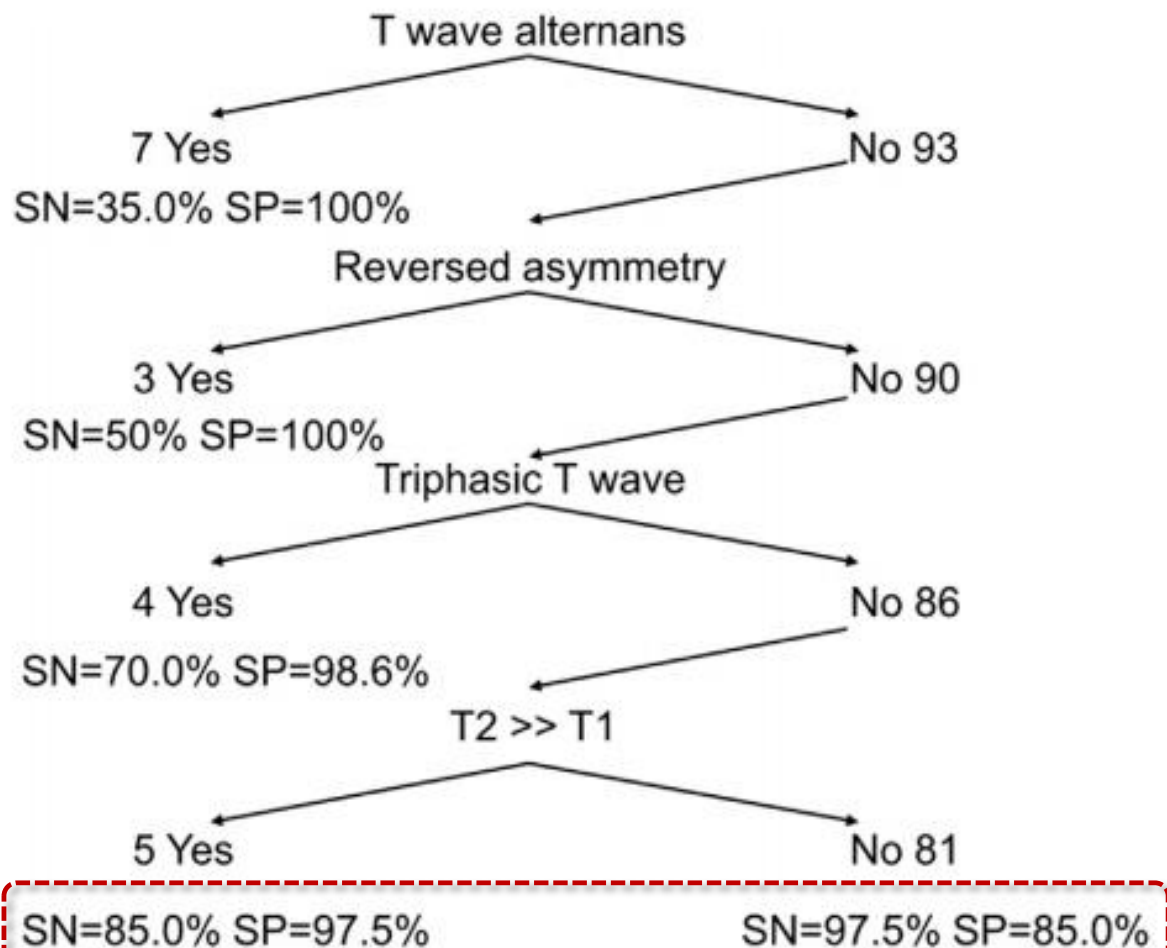
NPV = negative predictive value; PPV = positive predictive value; QTc = corrected QT interval.

LQT2-like T wave +
QT interval > 510ms
Tpeak-Tend > 85ms

Specificity 99%
PPV 94%

Specificity 97.3%
PPV 84.2%

Electrocardiographic predictors of bradycardia-induced torsades de pointes in patients with acquired atrioventricular block



Summary

- TdP is a polymorphic ventricular tachycardia caused by a long QT interval.
- **Prolongation of AP** (decrease in K^+ or increase in Na^+ or Ca^{++}) → **transmural dispersion** of repolarization → **EAD** (reactivation of L-type Ca^{++} or late I_{Na} or Na-Ca exchanger) → **PVC** when EAD reaching threshold → initiation of **TdP** → perpetuation (transmural reentry)

Predictors for TdP risk from ECG

- **QT prolongation**

→ heterogeneous dispersion of repolarization

- **PVC with long-short sequence, R-on-T**

- **Changes in QRS morphology and axis**

- **Changes in T wave morphology**

- T [peak] – T [end] interval
- Notched T wave (similar to LQT2) – genetic component (+)
- Complex (triphasic) T wave
- T wave alternans (macroscopic beat-to-beat alteration)
- Long QT with abnormal T-U wave

Thank you for your attention !

