

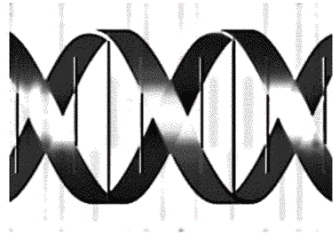
Ion channel mutations and disease: Insights from theoretical models

Genetic mutations in Na⁺ channels underlie clinical syndromes

- SCN5A (Cardiac) The Long-QT syndrome, Brugada Syndrome, Isolated cardiac conduction disorder
- SCN4A (Skeletal) hyperkalemic periodic paralysis, paramyotonia congenita and potassium-aggravated myotonias
- SCN1A (CNS) epilepsy, familial autism
- SCN2A (CNS) epilepsy
- SCN3A (CNS) familial autism
- SCN8A (CNS) epilepsy

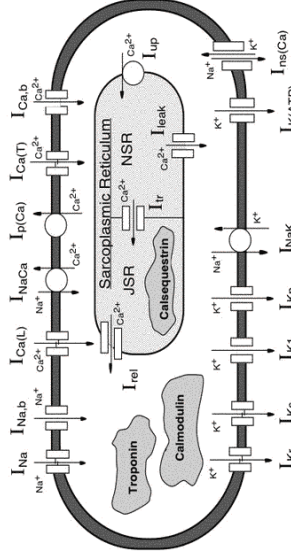
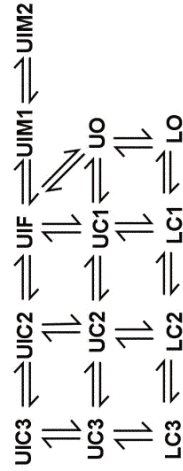
Integration:

1. genetic mutation

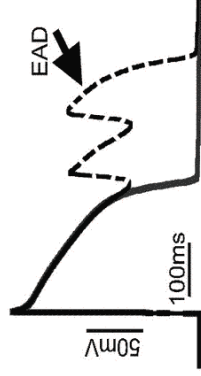


3p 21-24
LQT3, SCN5A

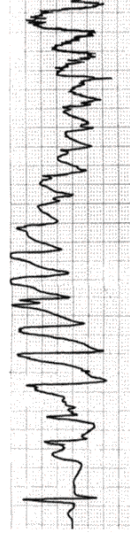
2. altered channel function



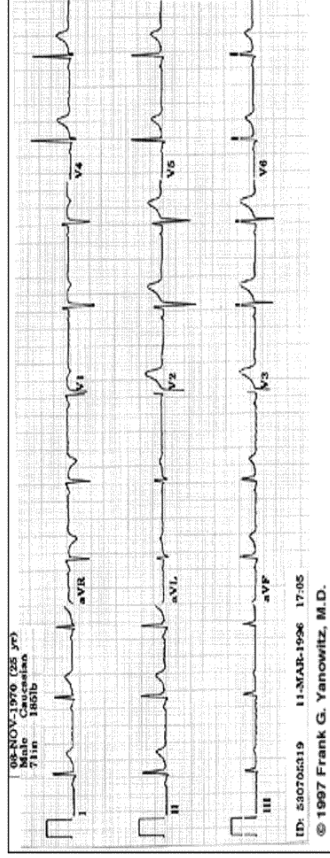
3. cardiac cell model



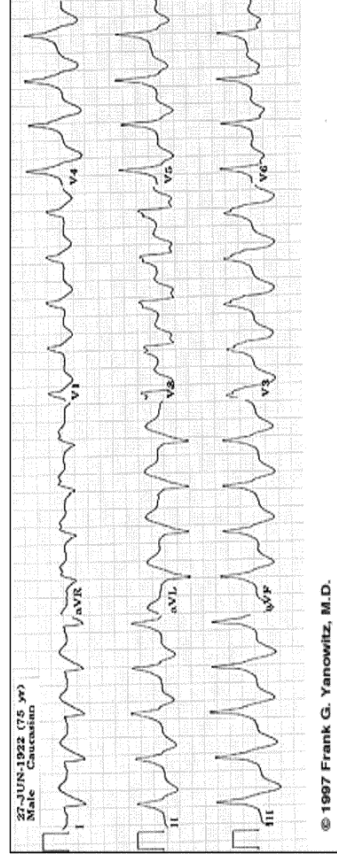
4. cardiac arrhythmia



Normal ECG

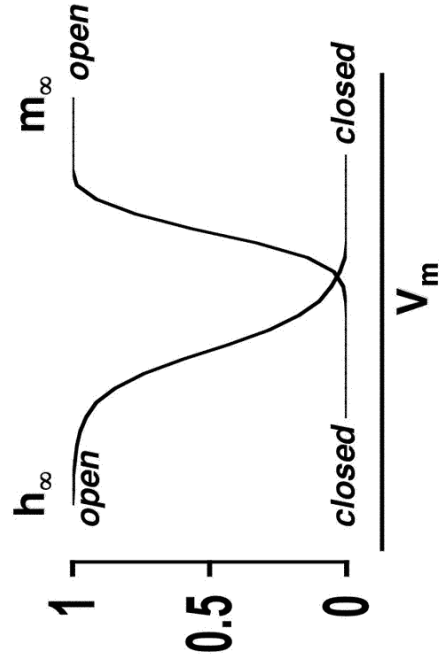


Ventricular tachycardia

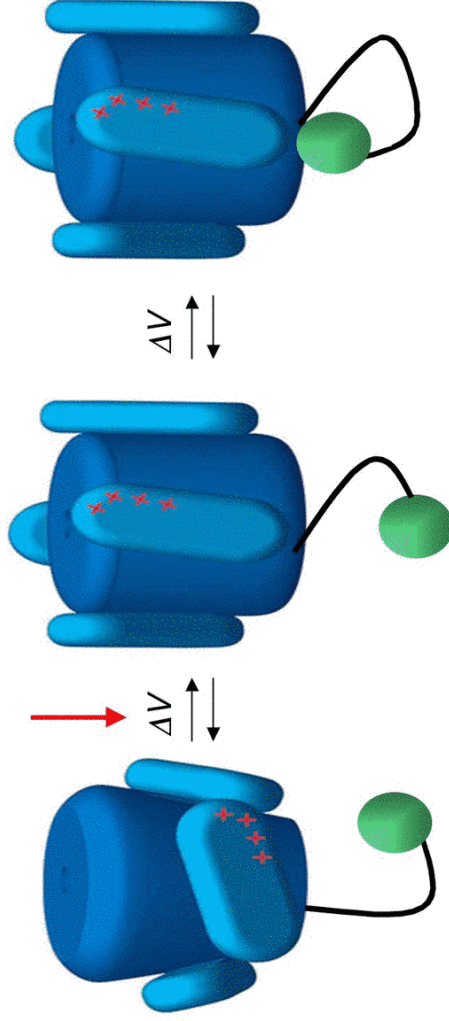
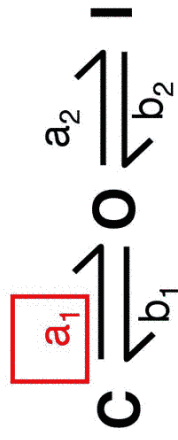


The Hodgkin-Huxley framework

$$I_{Na} = m^3 h^* g_{Na} * (E - E_{Na})$$



Markov models can incorporate mutations that affect discrete transitions



Computation of state probabilities



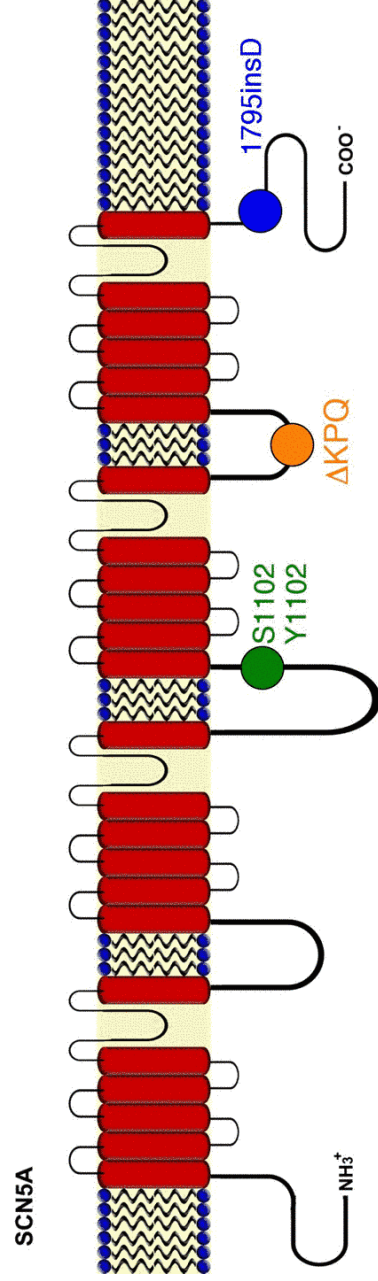
$$\frac{dP_i}{dt} = \sum_{j=1}^N [k_{ji} \cdot P_j(t, V_m)] - \sum_{j=1}^N [k_{ij} \cdot P_i(t, V_m)]$$

$$dC/dt = P(O) \cdot b_1 - P(C) \cdot a_1$$

$$dO/dt = P(C) \cdot b_1 + P(I) \cdot b_2 - (P(O) \cdot (a_2 + b_1))$$

$$dI/dt = 1 - P(O) - P(C)$$

Cardiac I_{Na}



ΔKPQ
The first mutation linked to Long-QT Syndrome

Three amino acid deletion in the channel inactivation gate.

1795insD*

Single amino acid insertion underlies

BOTH

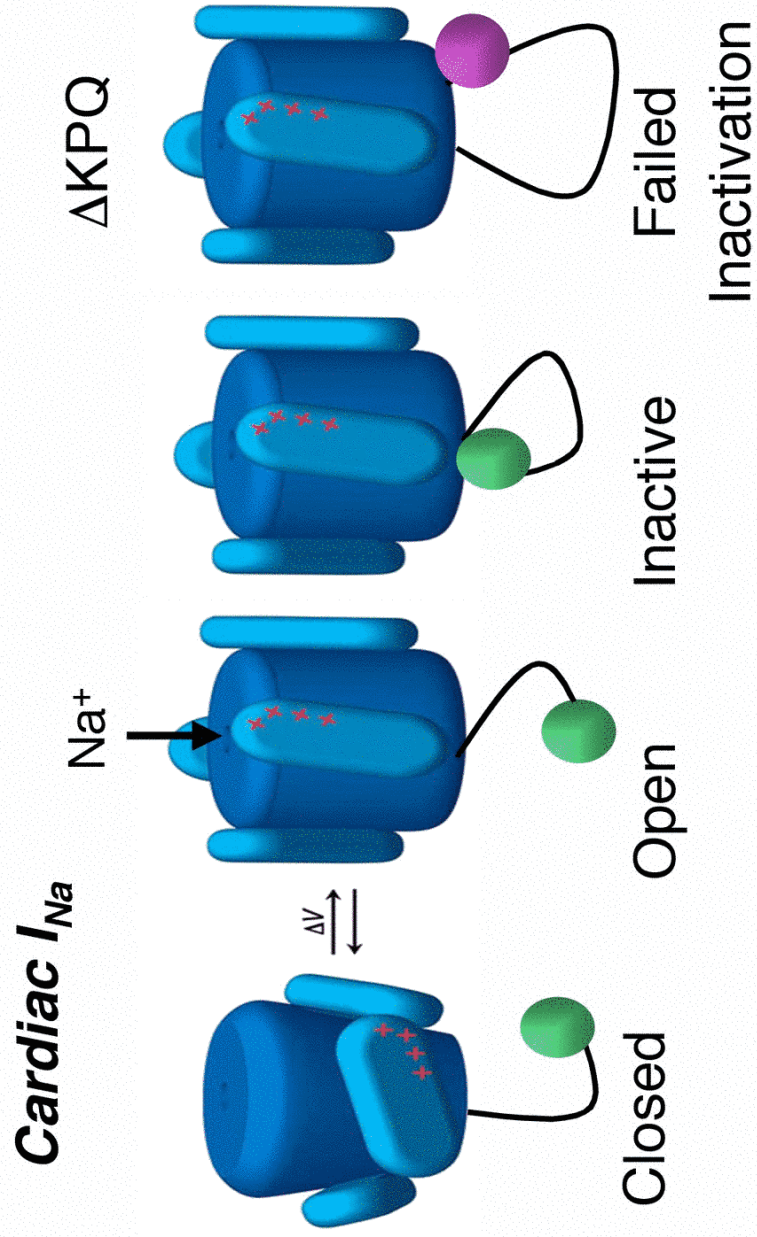
Long QT and Brugada Syndromes

S or Y 1102**

Genetic polymorphism in 13.2% of african americans

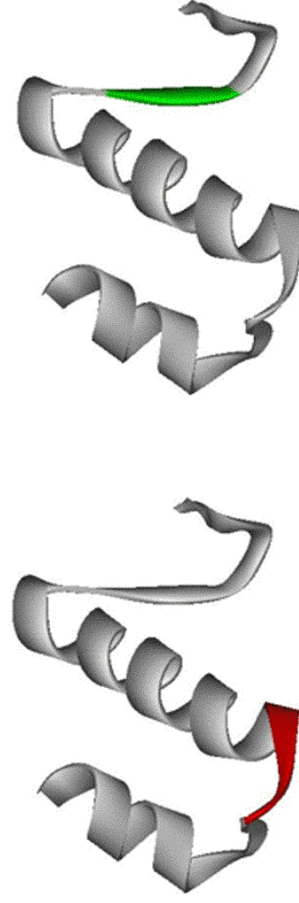
Predisposition to drug induced arrhythmia.

*Clancy and Rudy. 1999. Nature. 400. 566-569.



$\Delta K P Q$ channels may fail to inactivate

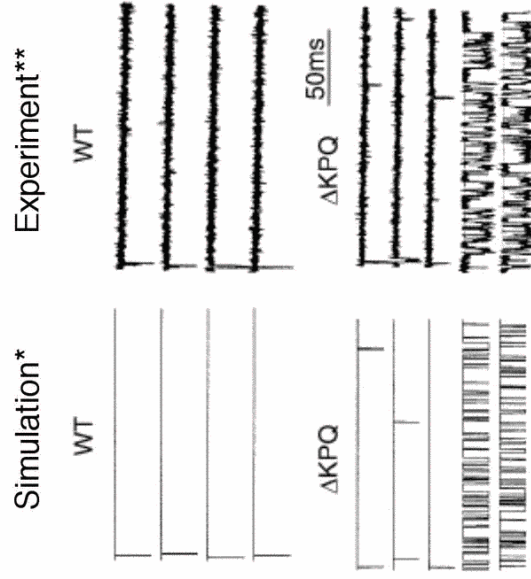
The $N a^{+}$ channel III-IV linker \rightarrow Fast inactivation



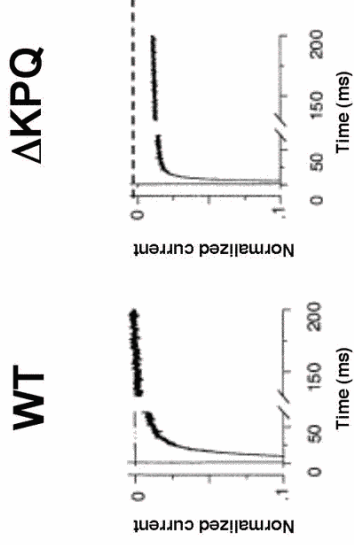
D N F N Q Q K K L G G Q D I F M T E E Q K K Y Y N A M K K L G S K K P Q K P I P R P L N K Y G F I F D

Kinetics of I_{Na}

Single channels



Whole cell***



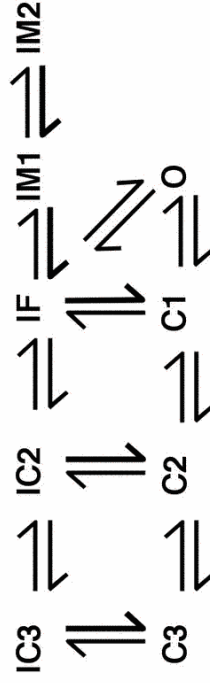
* Clancy and Rudy. 1999. Nature. 400. 566-569.

**Chandra, Starmer and Grant. 1998. AJP. H1643-H1654

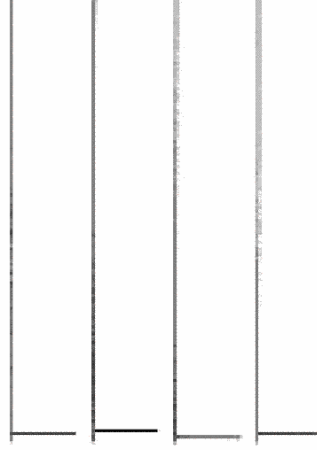
*** Bennett et al. 1995. Nature. 376. 683-685.

Modeling kinetics of I_{Na}

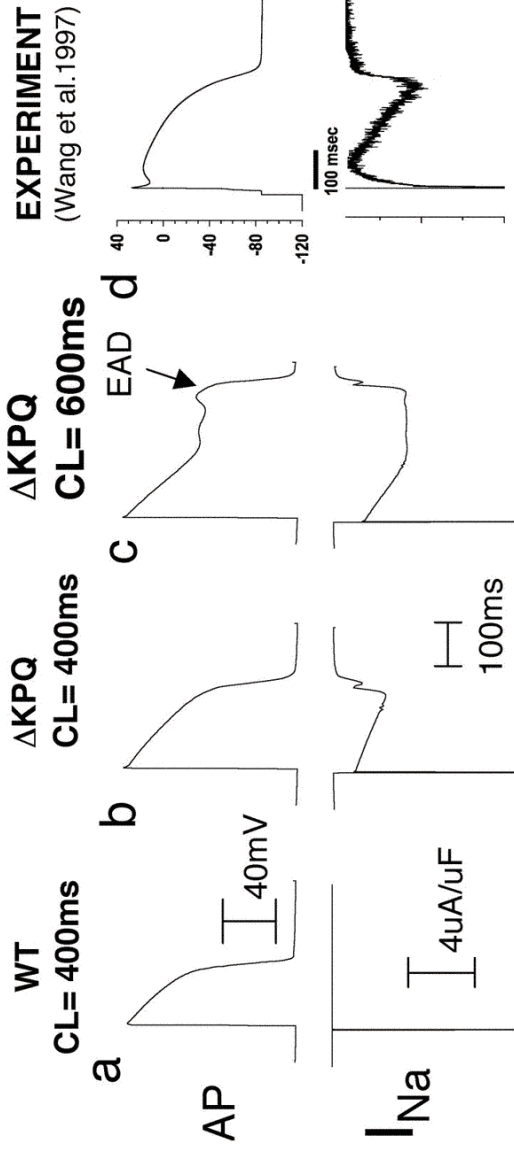
Markovian I_{Na}



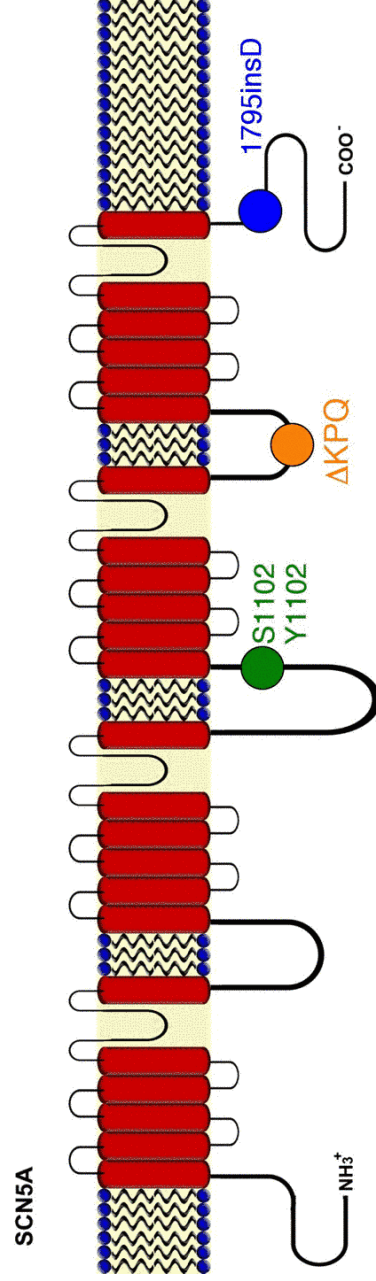
Simulated I_{Na} gating*



Δ KPQ Prolongs the Action Potential



Cardiac I_{Na}



Δ KPQ

Three amino acid deletion in the channel inactivation gate.



1795insD*

Single amino acid insertion underlies **BOTH** Long QT and Brugada Syndromes

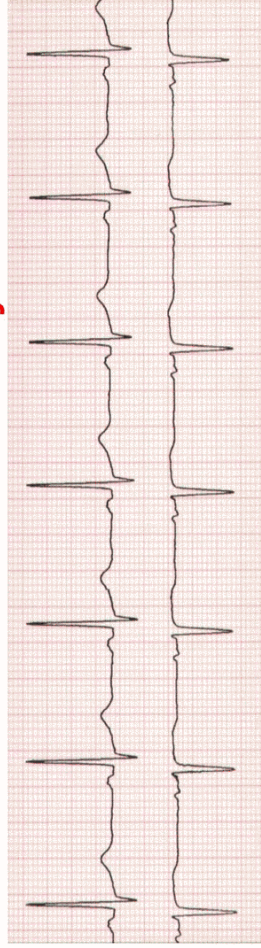


S or Y 1102**

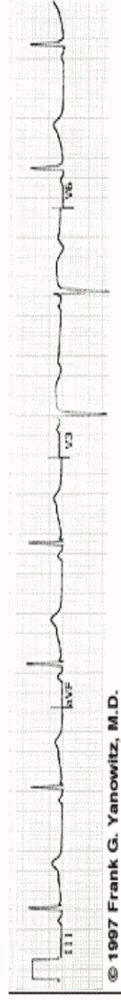
Genetic polymorphism in 13.2% of african americans
Predisposition to drug induced arrhythmia.

From Clancy and Rudy.
Circulation: 2002;105:1208-1213.

Normal Sinus Rhythm



Long QT interval

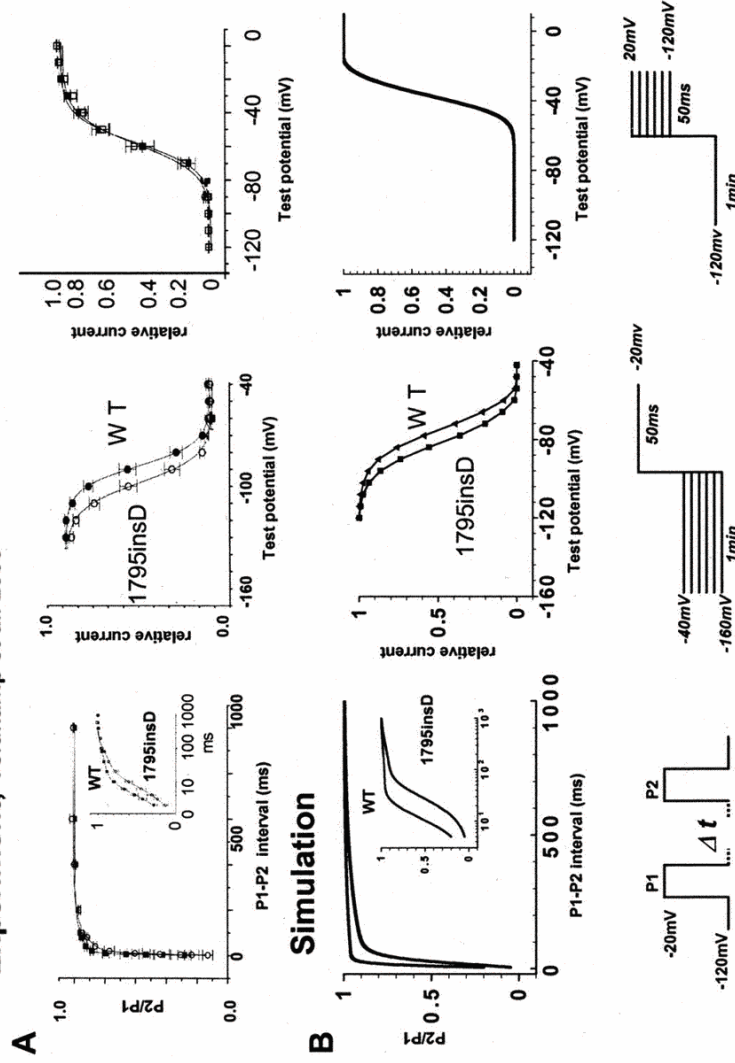


ST segment elevation



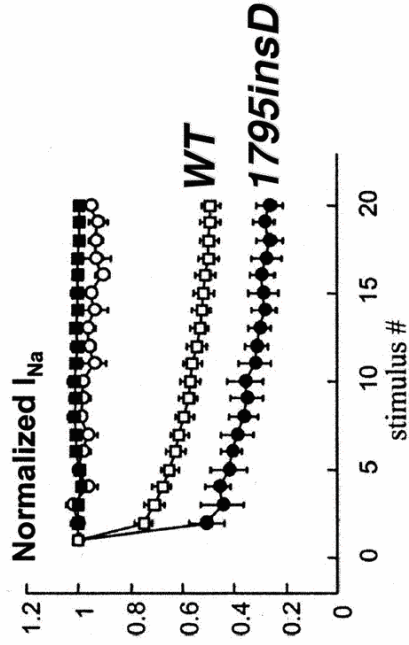
The 1795insD mutation slows recovery from inactivation and reduces channel availability

Experiment, Veldkamp et al. 2000

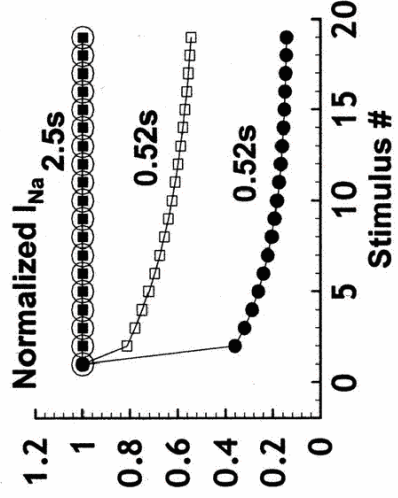


1795insD induced slower recovery from inactivation results in rate dependent current reduction

Experiment (Veldkamp et al.)

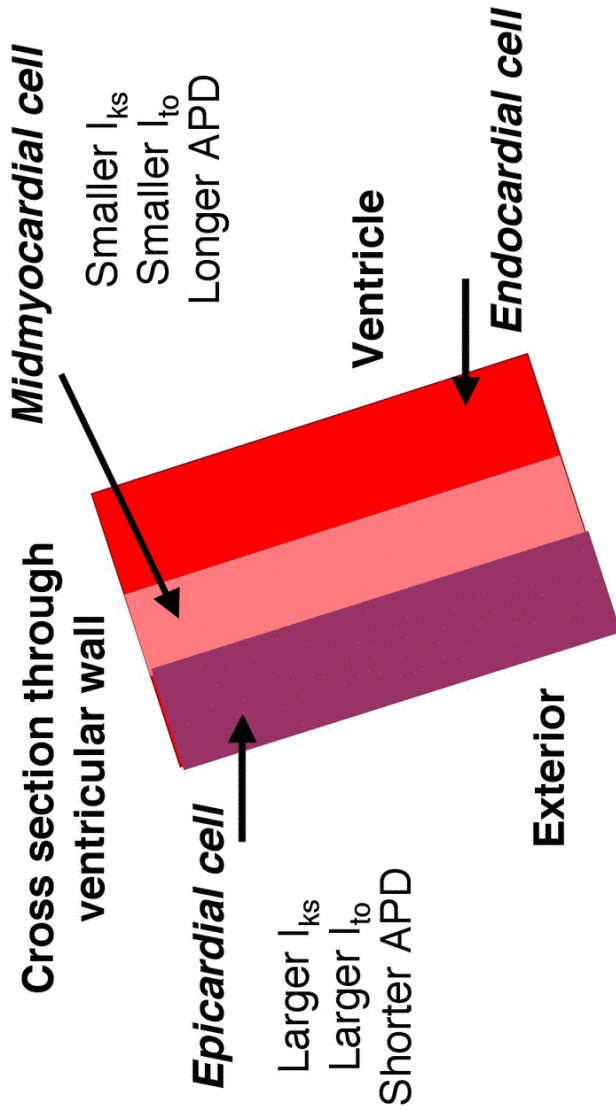


Simulation

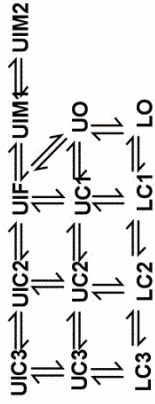


Epicardial vs. Midmyocardial cells

APD = Action Potential Duration



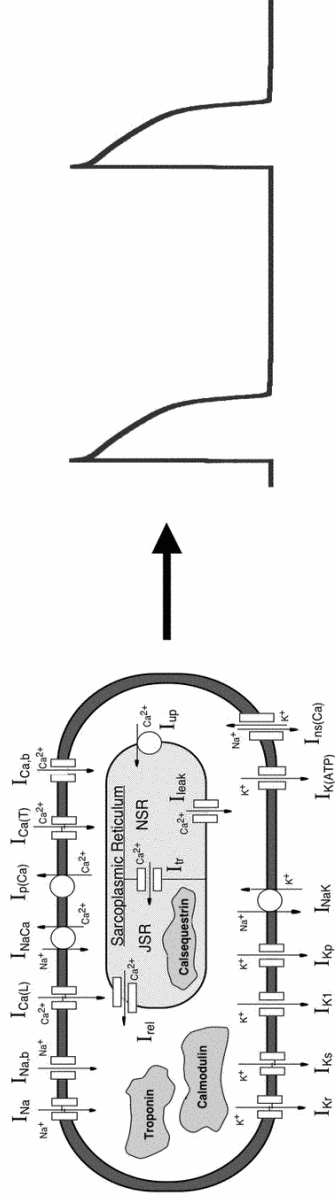
Markovian I_{Na}



Action potential simulations



Luo-Rudy AP model

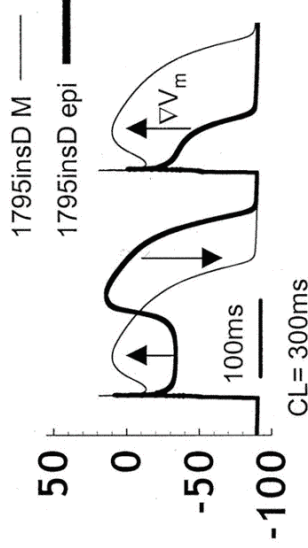


1795insD interaction with the heterogeneous myocardium

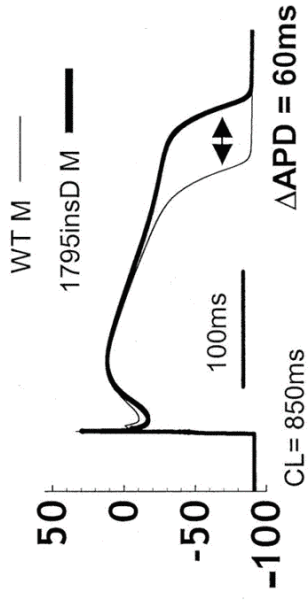
1795insD altered recovery kinetics underlies:

1795insD destabilizes inactivation (bursting channels) leading to:

A ST elevation

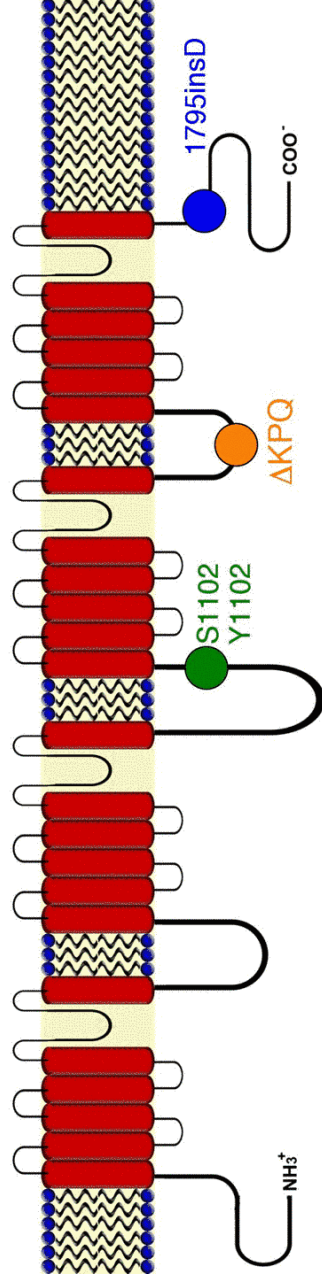


B QT prolongation



Cardiac I_{Na}

SCN5A



ΔKPQ

Three amino acid deletion in the channel inactivation gate.



1795insD*

Single amino acid insertion underlies

BOTH

Long QT and Brugada Syndromes



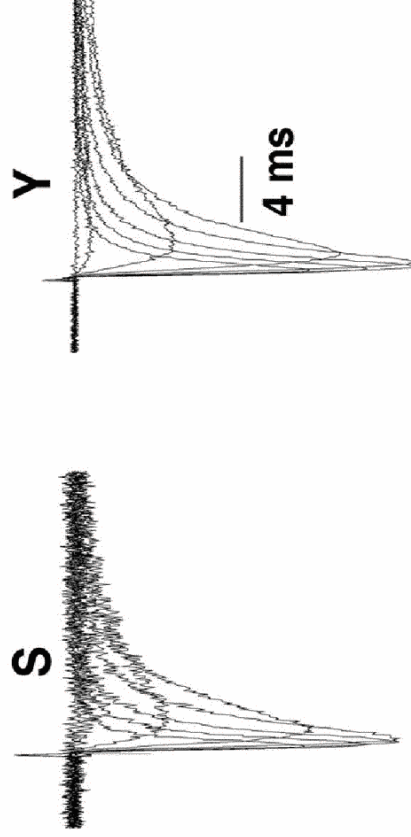
S or Y 1102**

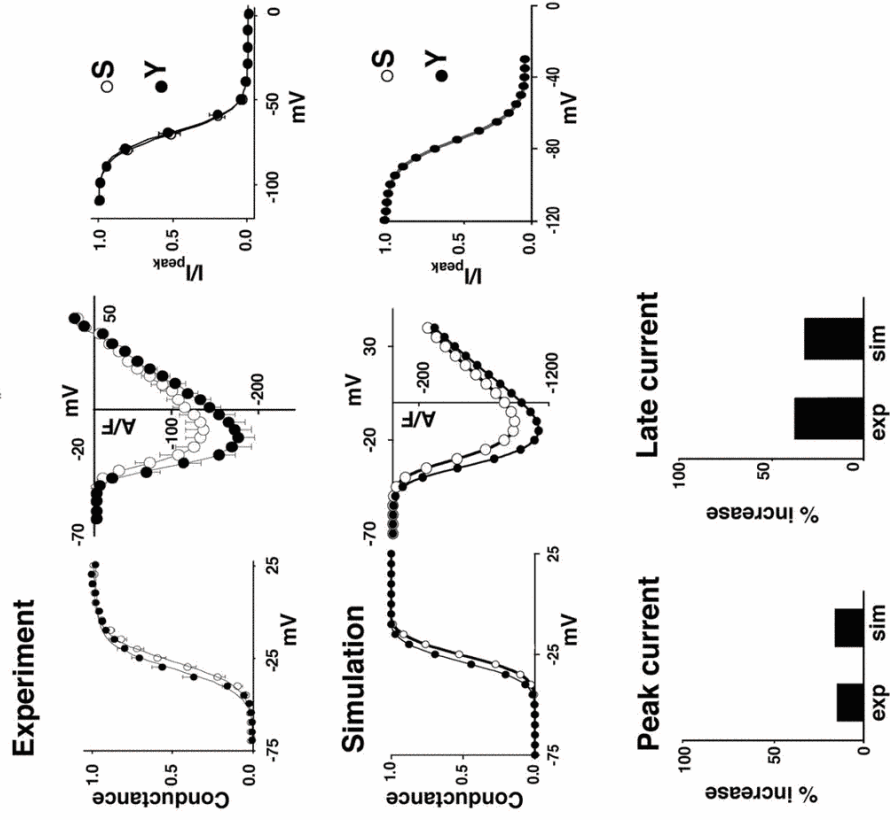
Genetic polymorphism in 13.2% of african americans
Predisposition to drug induced arrhythmia.

From Splawski, I. et al. 2002. Science. 297. 1333-36.

The Y1102 polymorphism has subtle effects on channel gating

Whole cell currents

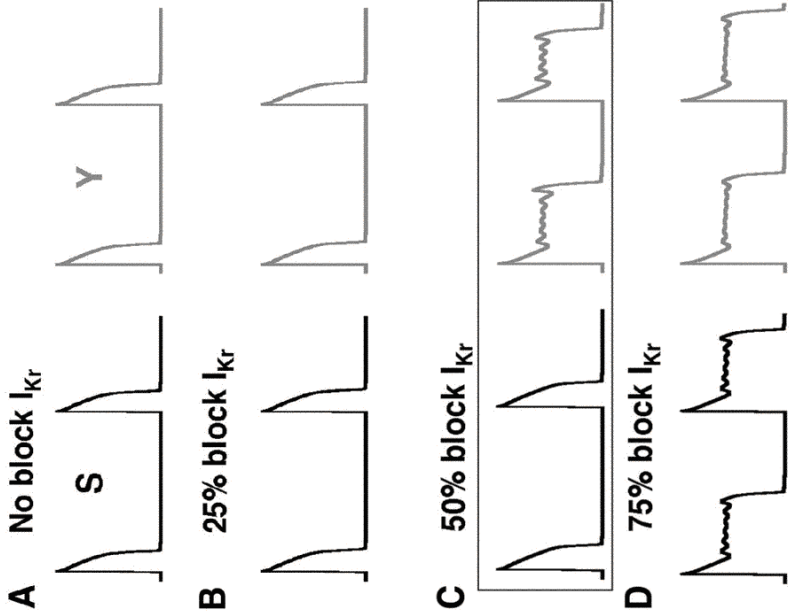




Y1102 causes a -5mV shift in activation gating.

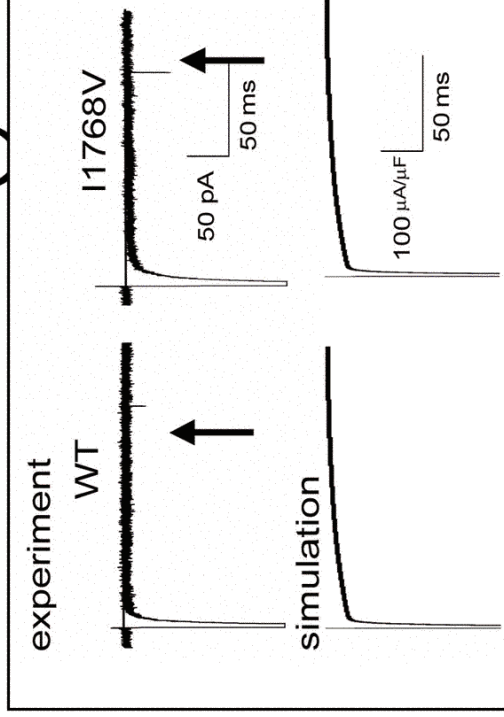
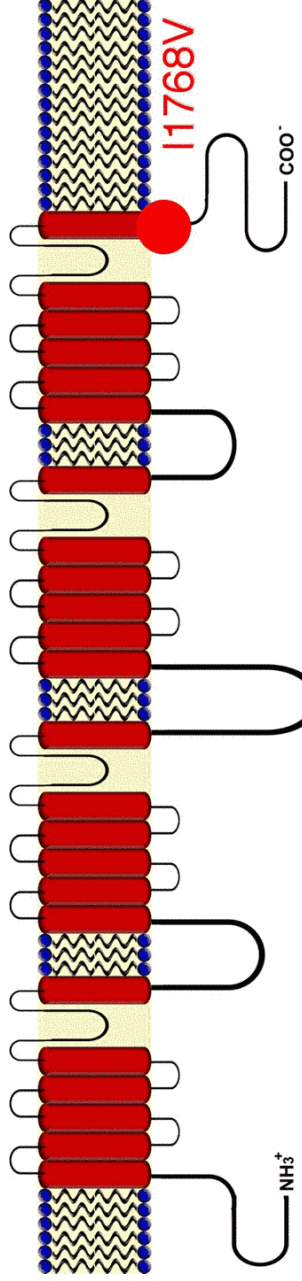
Consistent with experiments, the simulation predicts a 20% increase in peak current.

Y1102 facilitates development of triggered arrhythmias in the presence of K^+ channel block



The LQT-3 mutation I1768V is unlike the others

SCN5A

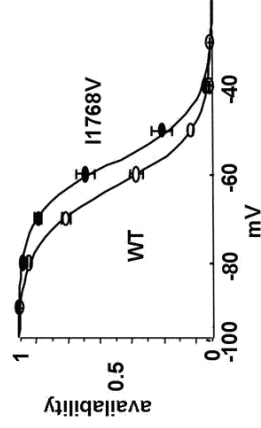
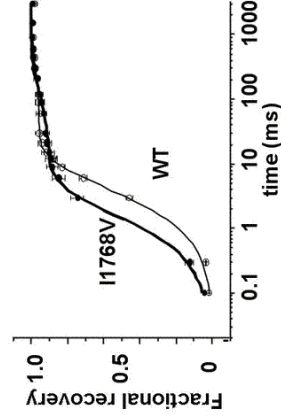


I1768V lacks persistent current

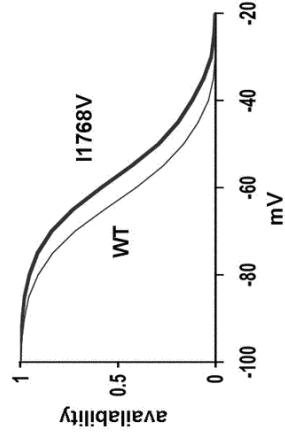
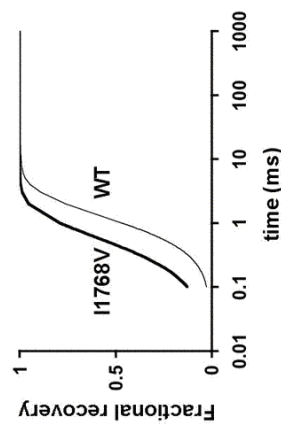
From Clancy, et al. 2003.
Circulation. 107(17):2233-7

I1768V alters channel recovery from inactivation

experiment

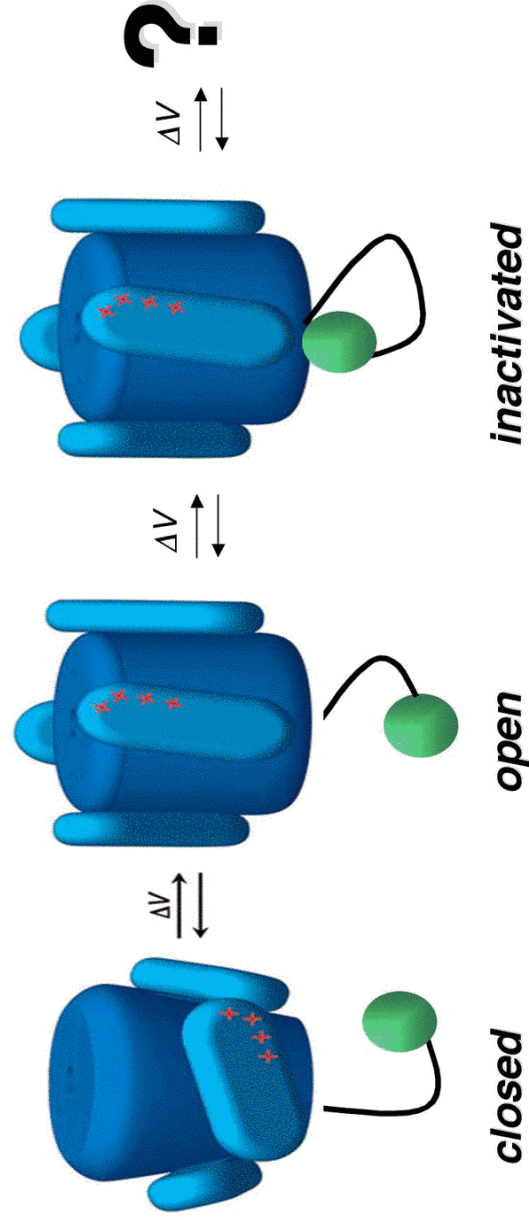


simulation



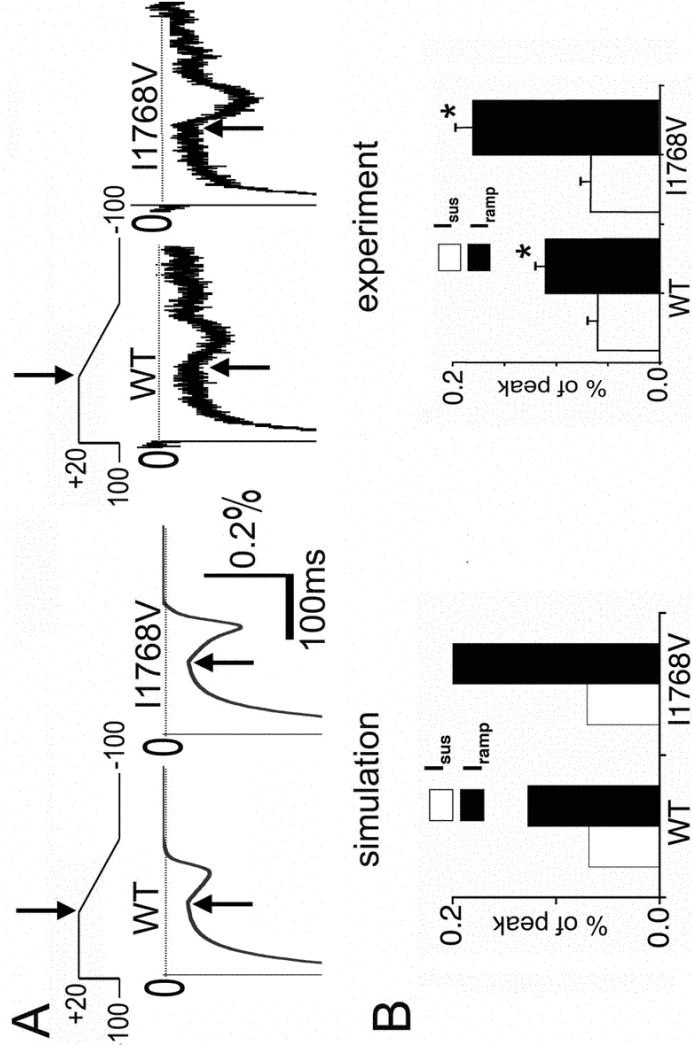
Question: Can we use theoretical techniques to guide experiments and elucidate the mechanism of I1768V induced Long-QT?

Exploration of **non-equilibrium gating** in cardiac Na⁺ channels

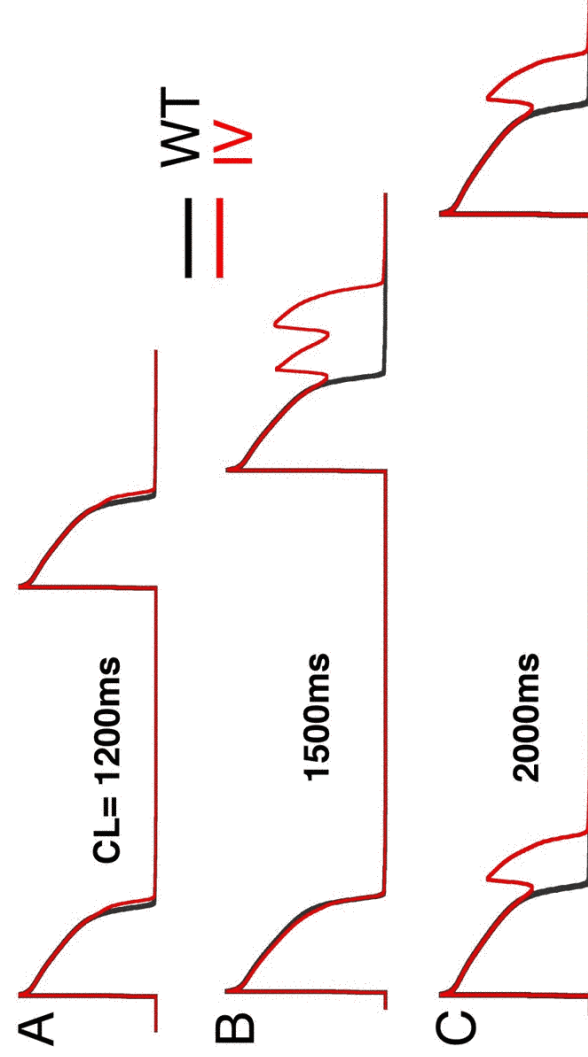


Does I1768V affect transitions out of the inactivation state?

The negative ramp protocol



1768V prolongs the action potential



Acknowledgements

Robert S. Kass (Columbia University)

Yoram Rudy (CWRU)

Michihiro Tateyama (Columbia University)

Igor Splawski (Harvard University)

Mark Keating (Harvard University)