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Study familial hypertrophic cardiomyopathy using patient-specific induced pluripotent stem cells

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Lab Meeting Journal Club 12th of October

Aims

Familial hypertrophic cardiomyopathy (HCM) is one the most common heart disorders, with gene mutations in the cardiac sarcomere. Studying HCM with patient-specific induced pluripotent stem-cell (iPSC)-derived cardiomyocytes (CMs) would benefit the understanding of HCM mechanism, as well as the development of personalized therapeutic strategies.

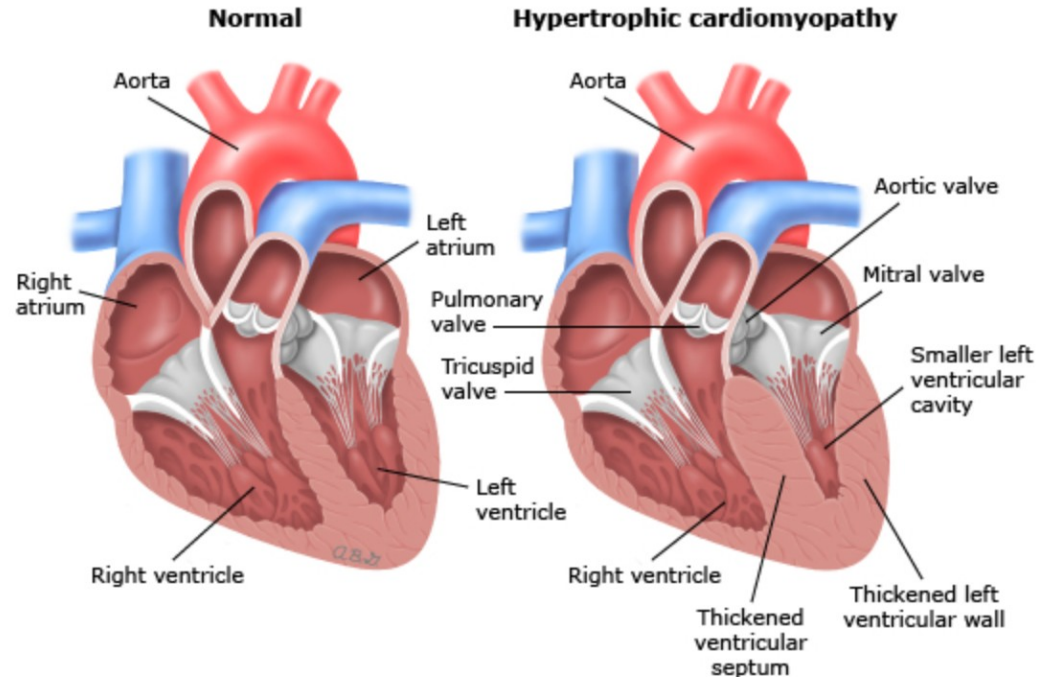
Hypertrophic Cardiomyopathy

Common symptoms:

- Chest pain
- Dyspnea
- Dizziness and fainting
- Fatigue

Complications:

- Sudden cardiac death
- Progressive heart failure
- Systolic dysfunction
- Stroke



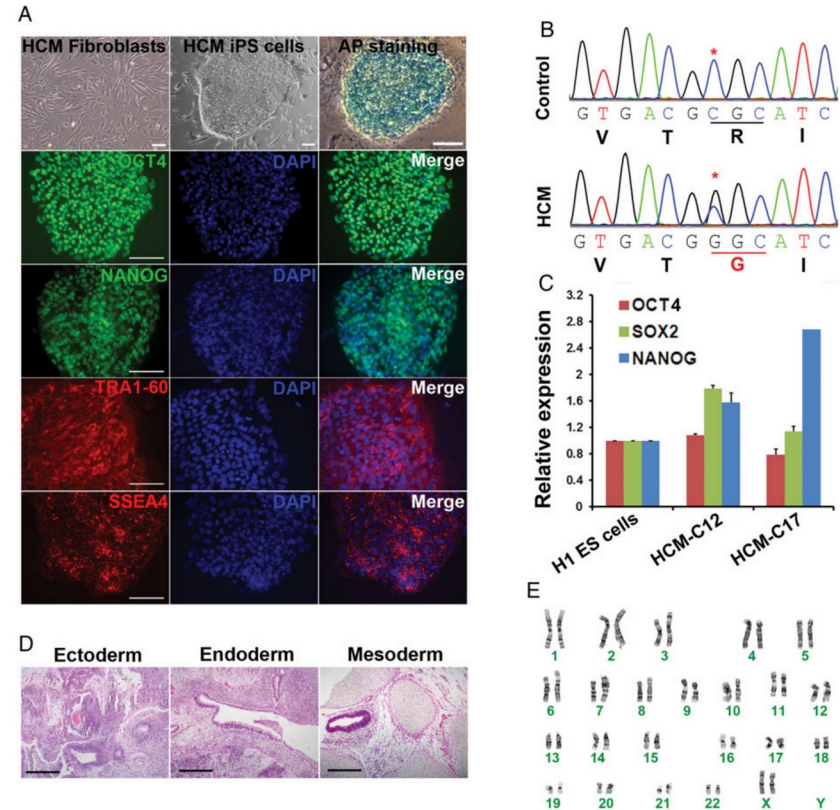
Genetic of HCM

Table 2 Genes associated with sarcomere hypertrophic cardiomyopathy

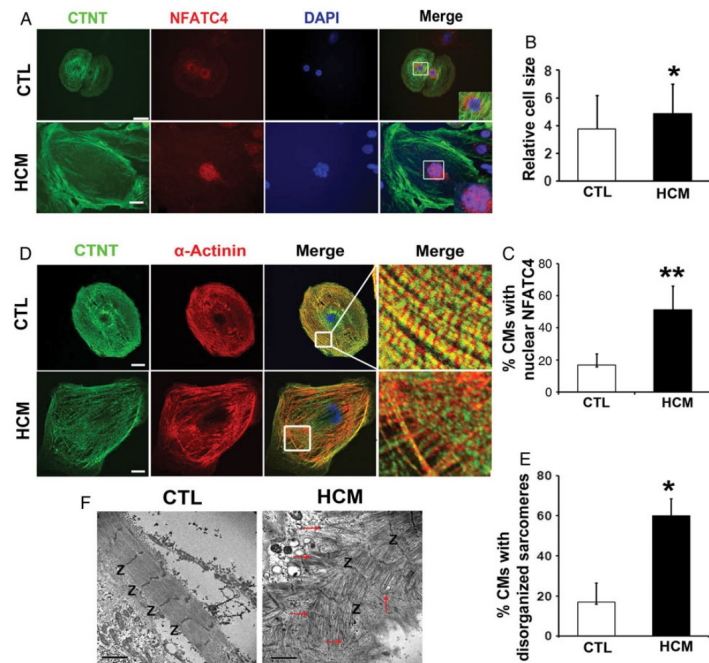
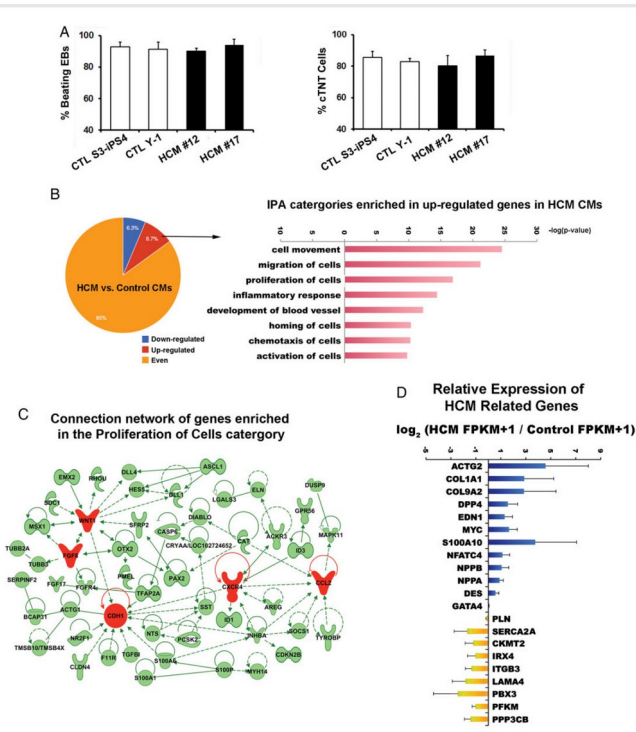
| Gene | Sarcomere protein | OMIM # | Frequency (%) |
|-------------------|-------------------------------------|--------|---------------|
| Thick myofilament | | | |
| <i>MYBPC3</i> | Myosin binding protein C | 600958 | ~40 |
| <i>MYH7</i> | β -myosin heavy chain | 160760 | ~40 |
| <i>MYL2</i> | Myosin light chain 2 | 160781 | <1 |
| <i>MYL3</i> | Myosin light chain 3 | 160790 | <1 |
| Thin myofilament | | | |
| <i>ACTC1</i> | Cardiac α -actin | 102540 | <1 |
| <i>TNNC1</i> | Cardiac troponin C | 191040 | <1 |
| <i>TNNI3</i> | Cardiac troponin I | 191044 | <5 |
| <i>TNNT2</i> | Cardiac troponin T2 | 191045 | ~10 |
| <i>TPM1</i> | α -tropomyosin | 191010 | <1 |
| Z-disc | | | |
| <i>ACTN2</i> | α -2 actinin | 102573 | <1 |
| <i>CSRP3</i> | Cysteine and glycine-rich protein 3 | 600824 | <1 |
| <i>MYOZ2</i> | Myozenin 2 | 605602 | <1 |
| <i>TCAP</i> | Telethonin | 604488 | <1 |
| <i>TTN</i> | Titin | 188840 | <1 |

HCM iPSC reprogramming and genotyping

Establishment and
characterization of HCM iPSCs



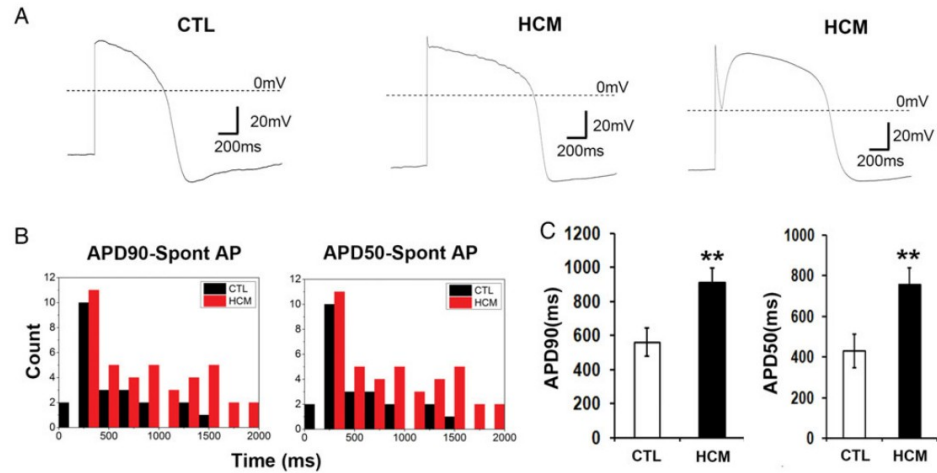
CM differentiation and genome-wide transcriptional profiling



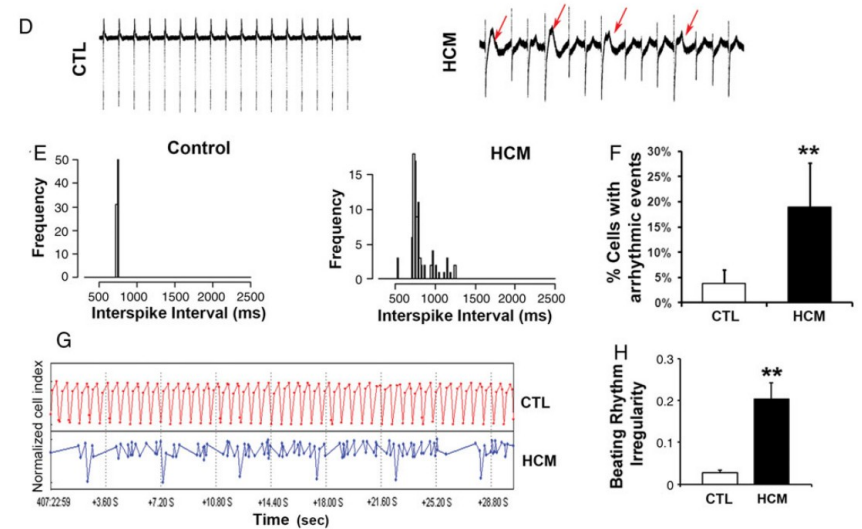
CM differentiation and gene expression profile of HCM iPSC-CMs

Phenotypic characterization of HCM iPSC-CMs

Electrophysiological analyses of HCM iPSC-CMs



Electrophysiological behaviour of single HCM iPSC-CMs



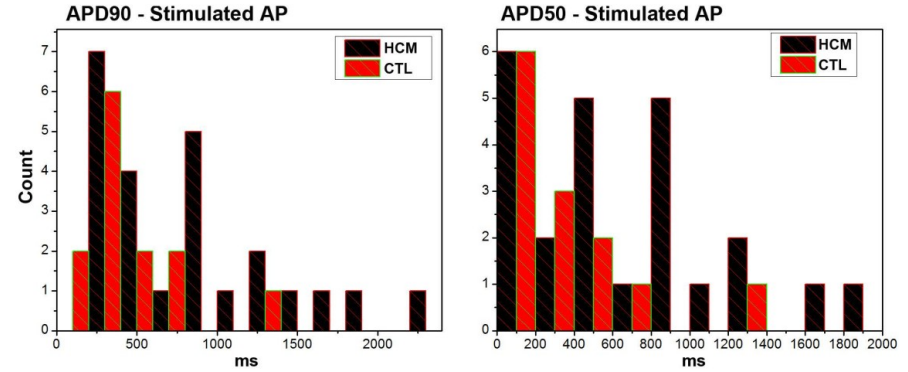
Electrophysiological behaviour of CM monolayers

Characterization of action-potentials Recording from the control and HCM IPSC-derived single cardiomyocytes.

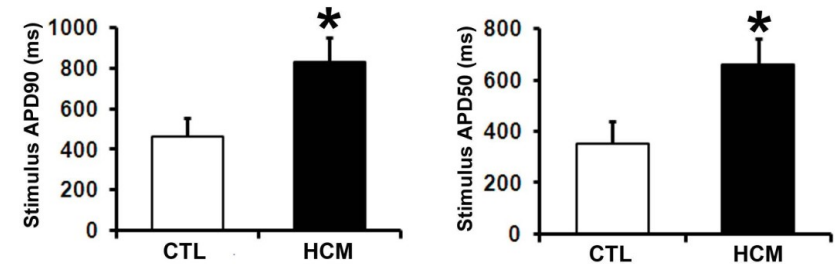
A

| | | Control | | HCM | |
|---|-------------------------------|----------|----------|----------|----------|
| | | Mean | SE | Mean | SE |
| Spontaneous APs | Minimum Diastolic Value (mV) | -58.441 | 2.212364 | -59.6188 | 1.905717 |
| | Mean Diastolic Potential (mV) | -49.199 | 2.167509 | -48.2268 | 1.894534 |
| | Amplitude | 79.67738 | 6.07583 | 86.24236 | 4.119675 |
| | Peak (mV) | 24.10327 | 2.688481 | 28.25734 | 2.620312 |
| | dV/dT max | 7.133617 | 0.433891 | 6.84151 | 0.284861 |
| | APD90 (ms) | 564.824 | 54.6295 | 711.7466 | 43.68447 |
| | APD50 (ms) | 450.2235 | 99.16871 | 595.9008 | 90.7935 |
| | V _m (mV) | -56.0027 | | -58.1559 | |
| Stimulated APs after a train of 4 beats at 0.5 Hz | Amplitude | 90.10721 | 4.353515 | 94.40086 | 2.337365 |
| | Peak (mV) | 34.10453 | 4.03087 | 36.23168 | 1.954102 |
| | dV/dT max | 78.99483 | 2.803166 | 86.70434 | 2.336736 |
| | APD90 (ms) | 560.4995 | 81.31584 | 910.3749 | 86.85604 |
| | APD50 (ms) | 429.3339 | 84.10154 | 756.3583 | 82.22055 |
| | Capitance | 65.99569 | 4.762481 | 66.55331 | 4.223107 |

B

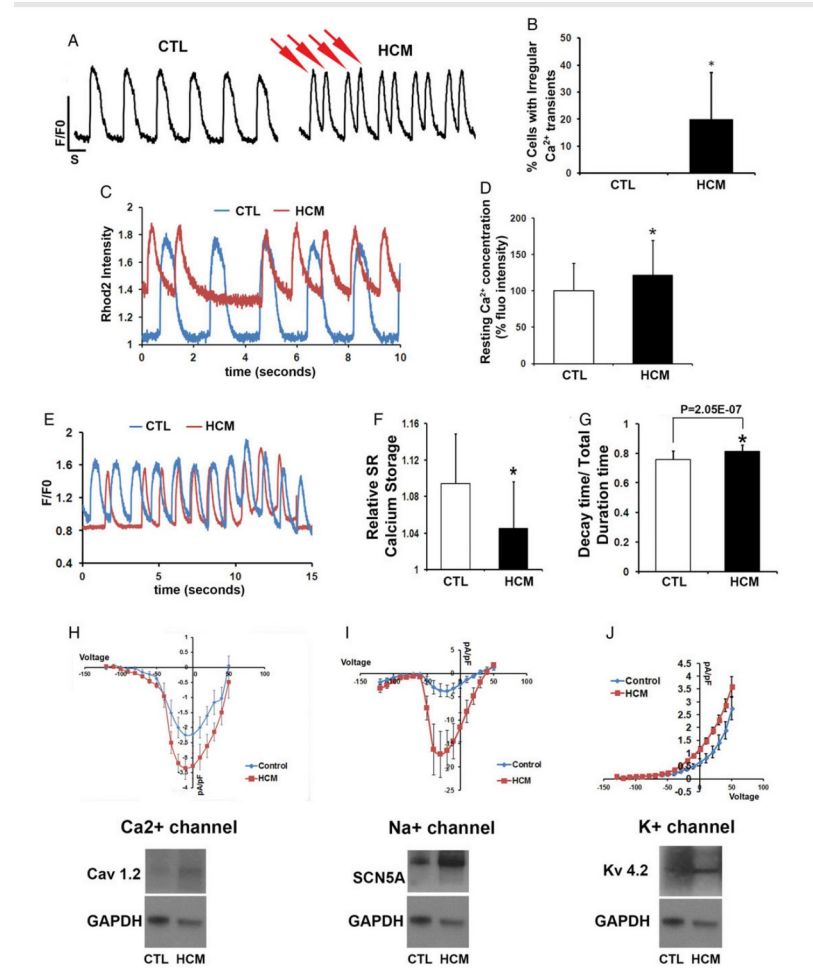


C

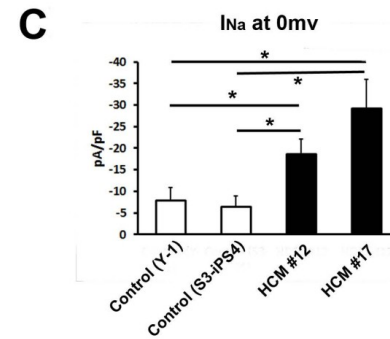
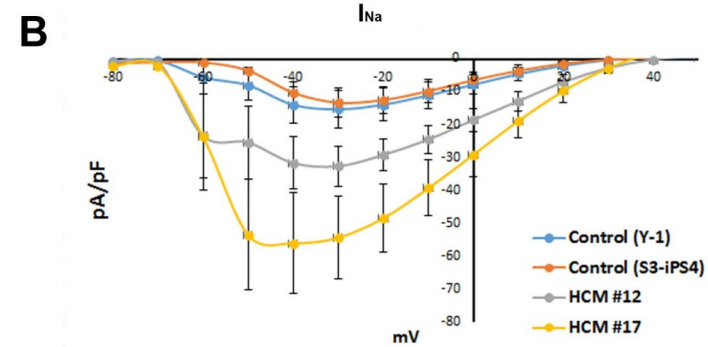
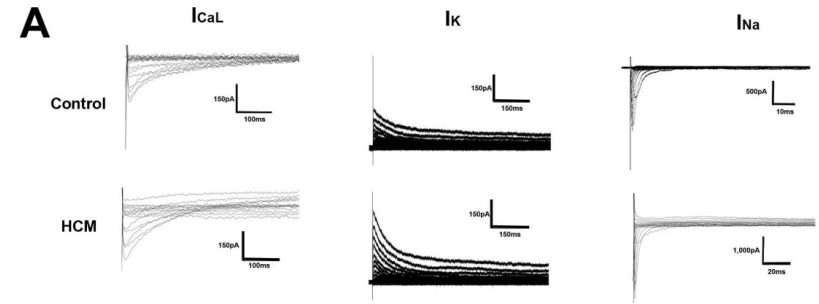


Calcium transient behaviour in HCM iPSC-CMs

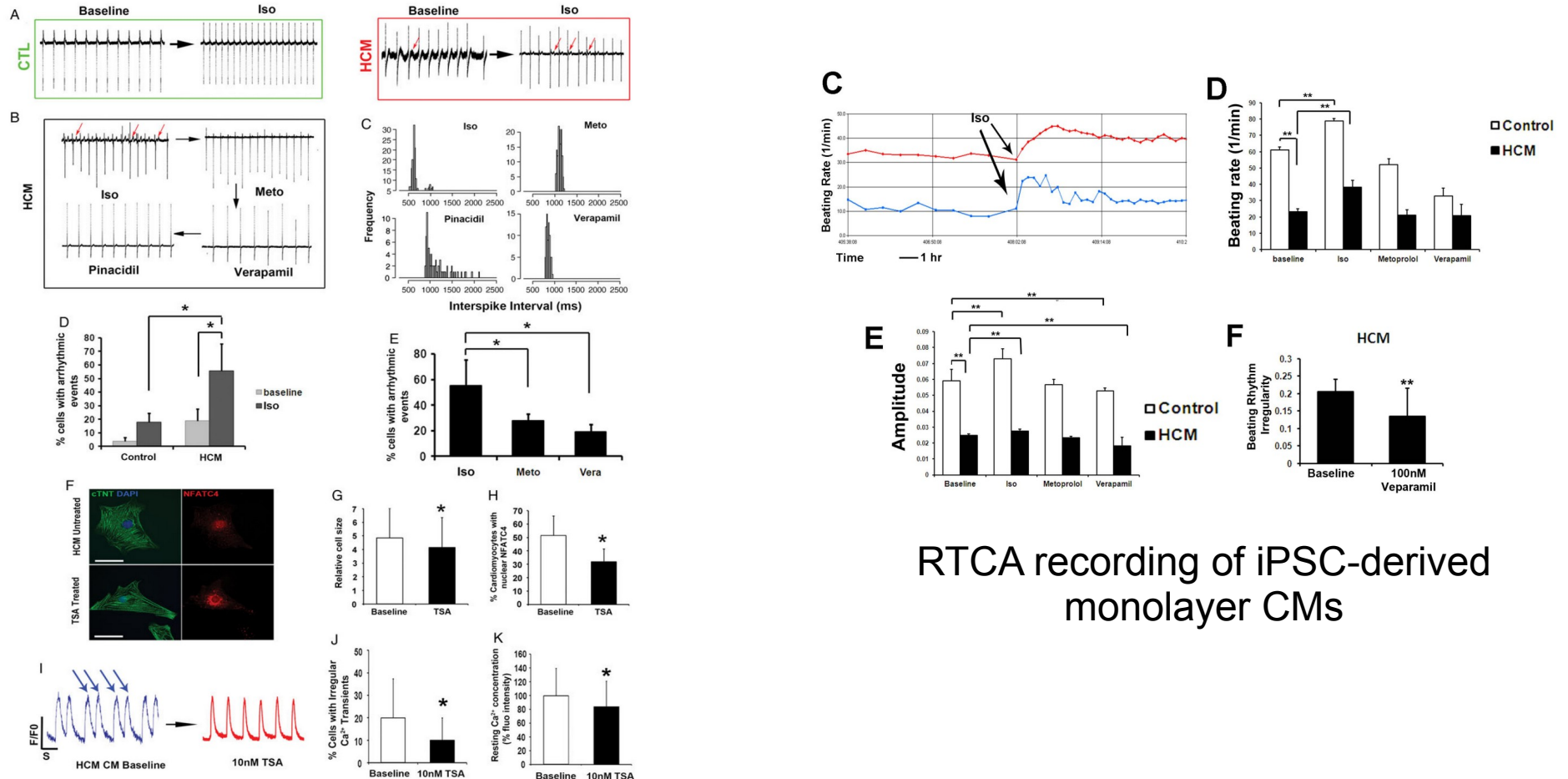
Analysis of calcium handling
and ion channels



Ion channel current changes in HCM IPSC cardiomyocytes



Pharmaceutical treatment of HCM iPSC-CMs



Research Article

Mutation-Specific Phenotypes in hiPSC-Derived Cardiomyocytes Carrying Either Myosin-Binding Protein C Or α -Tropomyosin Mutation for Hypertrophic Cardiomyopathy

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Aim: study the properties of HCM cardiomyocytes (CMs) derived from patient-specific human induced pluripotent stem cells (hiPSCs) carrying either MYBPC3-Gln1061X or TPM1-Asp175Asn mutation.

Action Potential Characteristics of WT and HCM hiPSC-Derived Cardiomyocytes

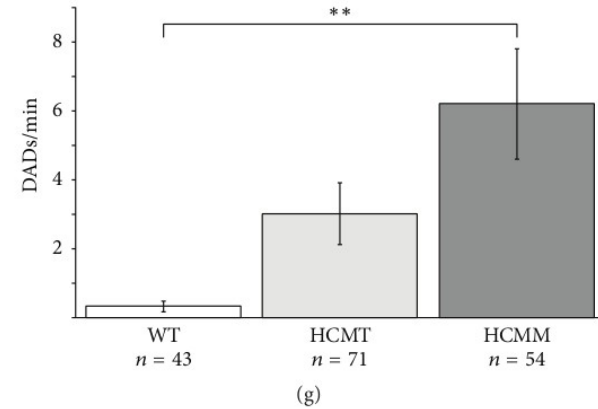
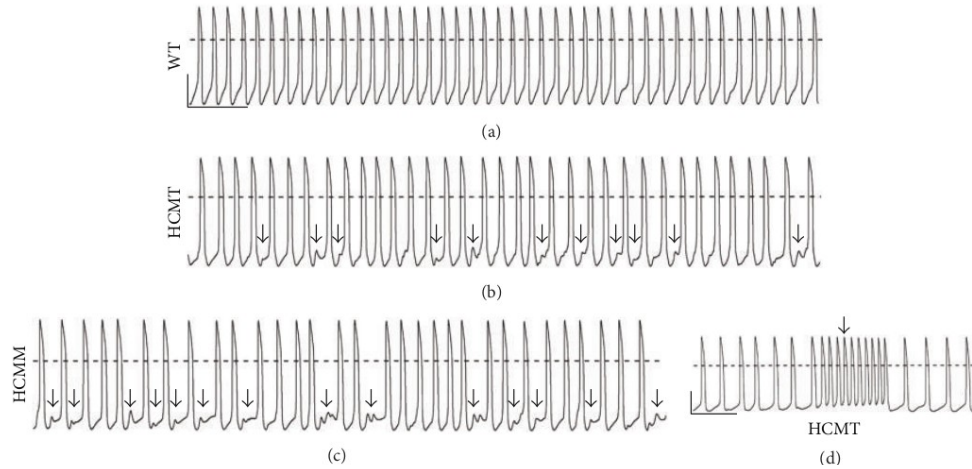


TABLE 3: AP properties of ventricular-like CMs derived from control hiPSC lines (WT) and from hiPSC lines carrying *TPM1-Asp175Asn* (HCMT) or *MYBPC3-Gln1061X* (HCMM) mutations. In the results, the data of each group is comprised from two separate cell lines.

| Group | n | Beating rate (BPM) | APD ₅₀ (ms) | APD ₉₀ (ms) | APA (mV) | MDP (mV) |
|-------|-----|-----------------------|---------------------------|---------------------------|-------------------|-----------------|
| WT | 43 | 58.1 ± 2.3 | 277.3 ± 13.0 | 323.6 ± 13.9 | 119.5 ± 1.1 | -76.8 ± 0.8 |
| HCMT | 71 | $48.4 \pm 1.5^{**}$ | $372.3 \pm 13.2^{**}$ | $433.1 \pm 14.0^{**}$ | 121.2 ± 1.1 | -75.8 ± 0.7 |
| HCMM | 54 | $47.1 \pm 1.8^{**}$ | $319.5 \pm 13.7^{\$}$ | $377.6 \pm 15.0^{*,\$}$ | $124.3 \pm 1.4^*$ | -77.9 ± 0.8 |

* HCMT or HCMM versus WT.

$\$$ HCMM versus HCMT.

$\$$ or * $p < 0.05$ and ** $p < 0.005$.