Original Articles

- Diminished PRRX1 Expression Is Associated With Increased Risk of Atrial Fibrillation and Shortening of the Cardiac Action Potential
- Genome-Wide Association Study to Find Modifiers for Tetralogy of Fallot in the 22q11.2 Deletion Syndrome Identifies Variants in the GPR88 Locus on 5q14.3
- New Blood Pressure–Associated Loci Identified in Meta-Analyses of 475,000 Individuals
- Transcriptome and Functional Profile of Cardiac Myocytes Is Influenced by Biological Sex
- Reduced Cardiac Calcineurin Expression Mimics Long-Term Hypoxia-Induced Heart Defects in Drosophila
- Body Height and Incident Risk of Venous Thromboembolism: A Cosibling Design
- Haritability of Mitral Regurgitation: Observations From the Framingham Heart Study and Swedish Population
- Validation of Polygenic Scores for QT Interval in Clinical Populations
- Prediction for Intravenous Immunoglobulin Resistance by Using Weighted Genetic Risk Score Identified From Genome-Wide Association Study in Kawasaki Disease

Editorial: One Size Does Not Fit All: Genetic Prediction of Kawasaki Disease Treatment Response in Diverse Populations
Impact of Selection Bias on Estimation of Subsequent Event Risk
Editorial: Biases in Genetic Association of Coronary Heart Disease Events May Be Less Likely Than Suspected: Here Is When to Check for Them
Predicting the Functional Impact of KCNQ1 Variants of Unknown Significance
Editorial: Machine Learning and Rare Variant Adjudication in Type 1 Long QT Syndrome
Characterization of a Human Induced Pluripotent Stem Cell–Derived Cardiomyocyte Model for the Study of Variant Pathogenicity: Validation of a KCNJ2 Mutation
Editorial: Commercially Available Human-Induced Pluripotent Stem Cell–Derived Cardiomyocytes: Another Piece in Our Tool Box, but Not a Swiss Army Knife Yet
α-Galactosidase A Genotype N215S Induces a Specific Cardiac Variant of Fabry Disease
Editorial: Clinical Characteristics of the GLA N215S Variant and Implications for the Diagnosis and Management of Nonclassic Fabry Disease
A Comparison of Whole Genome Sequencing to Multigene Panel Testing in Hypertrophic Cardiomyopathy Patients
Editorial: Hypertrophic Cardiomyopathy Gene Testing: Go Big?
Genomic Triangulation and Coverage Analysis in Whole-Exome Sequencing–Based Molecular Autopsies
Editorial: Genomic Triangulation in Sudden Unexplained Death in the Young: The Way to Go?
Care in Specialized Centers and Data Sharing Increase Agreement in Hypertrophic Cardiomyopathy Genetic Test Interpretation
Editorial: Pathogenicity of Hypertrophic Cardiomyopathy Variants: A Path Forward Together

Perspective

What Do We Really Think About Human Germline Genome Editing, and What Does It Mean for Medicine?

Clinical Genomic Cases

Heterozygous Null LDLR Mutation in a Familial Hypercholesterolemia Patient With an Atypical Presentation Because of Alcohol Abuse
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