



# INHERITED CARDIOMYOPATHIES Requisition

Ship to:

## Molecular Genetics Diagnostic Laboratory

Regional Genetics Program

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<http://www.cheo.on.ca/en/moleculargenetics>

Patient Name:	_____	_____	_____
	Last	First	Initial
Health Card Number:	_____		
DOB: (yy/mm/dd)	_____		
Address:	_____		
	_____		
Telephone:	_____		
Gender (circle one):	Male	Female	

### ALL SECTIONS MUST BE COMPLETED

Collection Date: \_\_\_\_\_

Collection Centre: \_\_\_\_\_

CHEO Pedigree Number: \_\_\_\_\_

### Sample Requirements

#### Blood

EDTA (lavender top) \_\_\_\_\_ mL room temp (3 ML for infants 1 year or less; otherwise 2 X 5 ML)

For any other sample types, please contact the laboratory directly.

### Health Care Provider Requesting Test

Name: \_\_\_\_\_

Registration Number: \_\_\_\_\_

Address: \_\_\_\_\_

Telephone: \_\_\_\_\_

FAX: \_\_\_\_\_

Copy to: Name: \_\_\_\_\_

Registration Number: \_\_\_\_\_

Address: \_\_\_\_\_

Telephone: \_\_\_\_\_

FAX: \_\_\_\_\_

### Test Requested (see next page for the clinical testing criteria and a list of the genes included in each panel)

- ARVC panel (7 genes)
- CPVT panel (2 genes)
- DCM panel (25 genes)
- HCM panel (19 genes)
- Pan Cardiomyopathy panel (45 genes) *Note: most appropriate for overlapping or atypical phenotypes. If a disease-specific panel is negative and you would like to reflex to this panel, please contact the laboratory.*
- Single gene testing (Specify Gene): \_\_\_\_\_
- Store DNA for future testing (DNA will be stored for 2 years then discarded; contact the laboratory directly if a longer storage term is required)

<input type="checkbox"/> Family Variant Specific Test <i>(Include a copy of the family member's genetic test report. A positive control is recommended if testing was performed in a different lab)</i>
Gene _____ Variant _____
Gene _____ Variant _____
Gene _____ Variant _____
Proband name: _____
Relationship to proband: _____

### Clinical Information

Clinical Diagnosis:  HCM  DCM  ARVC  CPVT  LVNC  RCM  CHD  Unknown  
 Unaffected  Other \_\_\_\_\_

#### Cardiovascular Features:

##### Left Ventricular Hypertrophy

Max. LV wall thickness: \_\_\_\_\_ (mm)

##### Ventricular Enlargement/Dilation

##### Reduced Ejection Fraction

##### Conduction Disease/Arrhythmia

Asymmetric  Concentric

Left  Right

Yes \_\_\_\_\_%  No

WPW  Ventricular Tachycardia

AV Block  Atrial Fibrillation

Other \_\_\_\_\_

#### Other Clinical Features:

Yes (specify below)  No  Unknown

#### Ethnicity: (be as specific as possible)

Ashkenazi Jewish  Black/African  East Asian  European  First Nations  French Canadian

Hispanic  Middle Eastern  South Asian  Other \_\_\_\_\_

Positive Family History  Yes (specify below)  No  Unknown

# INHERITED CARDIOMYOPATHIES TEST DETAILS

## Clinical criteria for genetic testing:

- 1) Patient has a clinical diagnosis of the relevant condition
- 2) Patient does not have a diagnosis of the relevant condition, but clinical suspicion is strong
- 3) Patient has a family history of a pathogenic or likely-pathogenic mutation; in this scenario, testing should consist of analysis for the familial mutation only

## Methodology of genetic testing:

- 1) Sequencing: analysis of coding sequences of the relevant genes and 10 base pairs immediately adjacent to each exon. This test is performed by oligonucleotide-based target capture (TruSight Cardiomyopathy Panel, Illumina) followed by next generation sequencing using the MiSeq instrument (Illumina). Additional Sanger sequencing is performed for relevant regions that have insufficient coverage, and to confirm all clinically significant variants and variants of unknown significance.
- 2) MLPA: to detect large genomic deletions and duplications, multiplex ligation-dependent probe amplification (MLPA) is performed for the relevant genes.

## Arrhythmogenic Right Ventricular Cardiomyopathy (ARVC) Panel

Genes included in panel: *DSC2*, *DSG2*, *DSP*, *JUP*, *PKP2*, *RYR2*, and *TMEM43*

Analysis includes sequencing as described above, and MLPA of *DSC2*, *DSG2*, *DSP* and *PKP2*.

## Catecholaminergic Polymorphic Ventricular Tachycardia (CPVT) Panel

Genes included in panel: *CASQ2* and *RYR2*

Analysis includes sequencing as described above.

## Hypertrophic Cardiomyopathy (HCM) Panel

Genes included in panel: *ACTC1*, *ACTN2*, *CAV3*, *CSRP3*, *GLA*, *LAMP2*, *MYBPC3*, *MYH7*, *MYL2*, *MYL3*, *MYOZ2*, *NEXN*, *PLN*, *PRKAG2*, *TNNC1*, *TNNI3*, *TNNT2*, *TPM1*, and *TTR*

Analysis includes sequencing as described above, and MLPA of *MYH7*, *MYBPC3* and *TNNT2*.

## Dilated Cardiomyopathy (DCM) Panel

Genes included in panel: *ABCC9*, *ACTC1*, *ACTN2*, *CSRP3*, *CTF1*, *DES*, *EMD*, *LAMP2*, *LDB3*, *LMNA*, *MYBPC3*, *MYH6*, *MYH7*, *NEXN*, *PLN*, *RBM20*, *SGCD*, *TAZ*, *TCAP*, *TNNC1*, *TNNI3*, *TNNT2*, *TPM1*, *TTN*, and *VCL*

Analysis includes sequencing as described above, and MLPA of *MYH7*, *MYBPC3* and *TNNT2*.

## Pan Cardiomyopathy Panel

Genes included in panel: *ABCC9*, *ACTC1*, *ACTN2*, *ANKRD1*, *CASQ2*, *CAV3*, *CRYAB*, *CSRP3*, *CTF1*, *DES*, *DSC2*, *DSG2*, *DSP*, *EMD*, *FHL2*, *GLA*, *JUP*, *LAMA4*, *LAMP2*, *LDB3*, *LMNA*, *MYBPC3*, *MYH6*, *MYH7*, *MYL2*, *MYL3*, *MYLK2*, *MYOZ2*, *NEXN*, *PKP2*, *PLN*, *PRKAG2*, *RBM20*, *RYR2*, *SGCD*, *TAZ*, *TCAP*, *TMEM43*, *TNNC1*, *TNNI3*, *TNNT2*, *TPM1*, *TTN*, *TTR*, and *VCL*

Analysis includes sequencing as described above, and MLPA of *DSC2*, *DSG2*, *DSP*, *MYH7*, *MYBPC3*, *PKP2* and *TNNT2*.

## Frequently Asked Questions about the Pan Cardiomyopathy Panel:

- 1) When is the pan cardiomyopathy panel appropriate to order (as opposed to a smaller disease-specific panel)?
  - When the patient has cardiomyopathy or strong clinical suspicion of cardiomyopathy, with features overlapping or atypical of conventional disease categories.
  - It is important to note that the pan cardiomyopathy panel is not a comprehensive arrhythmia panel. If the primary etiology is felt to be more likely a channelopathy, a comprehensive arrhythmia panel may be better suited. For instance, if a patient has VF arrest at 40 years or less, without a clear etiology identified, the pan cardiomyopathy panel is likely not the most appropriate first step.
- 2) Why wouldn't I always order the pan cardiomyopathy panel?
  - Because of the large number of genes analysed, the pan cardiomyopathy panel is likely to yield a higher number of variants of unclear clinical significance than the disease-specific panels. These variants can be difficult for the ordering provider and the patient to interpret.
  - If the patient's disease is consistent with an established cardiomyopathy, an expanded panel is expected to add only limited additional sensitivity [Alfares et al (2015) Genet Med, PMID 25611685].