



# NON-SYNDROMIC CARDIOMYOPATHIES Requisition

Ship to:  
**Genetics Diagnostic Laboratory**

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<https://www.cheo.on.ca/en/clinics-services-programs/genetics-diagnostic-laboratory.aspx>

## ALL SECTIONS MUST BE COMPLETED

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

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

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

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

## Sample Requirements

### Blood

Blood 2x 6 mL EDTA  Blood 2x 3 mL EDTA (child)  Blood 3 mL EDTA (infant ≤1 year)

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; see page 2 for details)
- Pan Cardiomyopathy panel (30 genes; see page 2 for details) **Note: most appropriate for DCM, or overlapping or atypical phenotypes.**
- Single gene testing (Specify Gene): \_\_\_\_\_
- Store DNA for future testing (DNA will be stored for 2 years then discarded)

Family Variant Specific Test  
(Include a copy of the family member's genetic test report. A positive control is recommended if testing was performed in a different lab)

Gene(s) \_\_\_\_\_

Variant(s) \_\_\_\_\_

Proband name: \_\_\_\_\_

Proband date of birth: \_\_\_\_\_

Relationship to proband: \_\_\_\_\_

Note: there is different requisition for hypertrophic cardiomyopathy genetic testing; it is available at <https://www.cheo.on.ca/en/clinics-services-programs/requisitions-and-forms.aspx>

## Clinical Information

**Clinical Diagnosis:**  HCM (Age of dx:\_\_\_\_)  DCM (Age of dx:\_\_\_\_)  ARVC (Age of dx:\_\_\_\_)  LVNC (Age of dx:\_\_\_\_)  
 Sudden cardiac arrest < 50 years old  Other: \_\_\_\_\_

### Cardiovascular Features:

<b>Hypertension</b> (treated with medication) <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown	<b>Ejection Fraction &lt;40%</b> <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown
<b>Pacemaker/ implantable defibrillator</b> <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown	<b>Peripartum onset</b> <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown
<b>Ventricular tachycardia</b> <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown	

**Ethnicity** (be as specific as possible; this is important as the frequency of rare DNA changes can vary between ethnic backgrounds):

- Ashkenazi Jewish  Black/African  East Asian  European  First Nations  French Canadian
- Hispanic  Middle Eastern  South Asian  Other \_\_\_\_\_

**Positive Family History (1<sup>st</sup> and 2<sup>nd</sup> degree relatives only)**  Yes (specify below)  No  Unknown  
 HCM (Age of dx:\_\_\_\_)  DCM (Age of dx:\_\_\_\_)  ARVC (Age of dx:\_\_\_\_)  LVNC (Age of dx:\_\_\_\_)  
 Sudden cardiac arrest/death < 50 years old  Cardiac transplant

Other: \_\_\_\_\_

## NON-SYNDROMIC CARDIOMYOPATHIES TEST DETAILS

### Methodology of genetic testing:

- 1) Sequencing: next-generation sequencing analysis of coding sequences of the relevant genes and 10 base pairs immediately adjacent to each exon. In addition, several deep intronic regions are analyzed for the presence of specific clinically relevant variants.
- 2) MLPA: to detect large genomic deletions and duplications, multiplex ligation-dependent probe amplification (MLPA) is performed for certain genes.

### Arrhythmogenic Right Ventricular Cardiomyopathy (ARVC) Panel

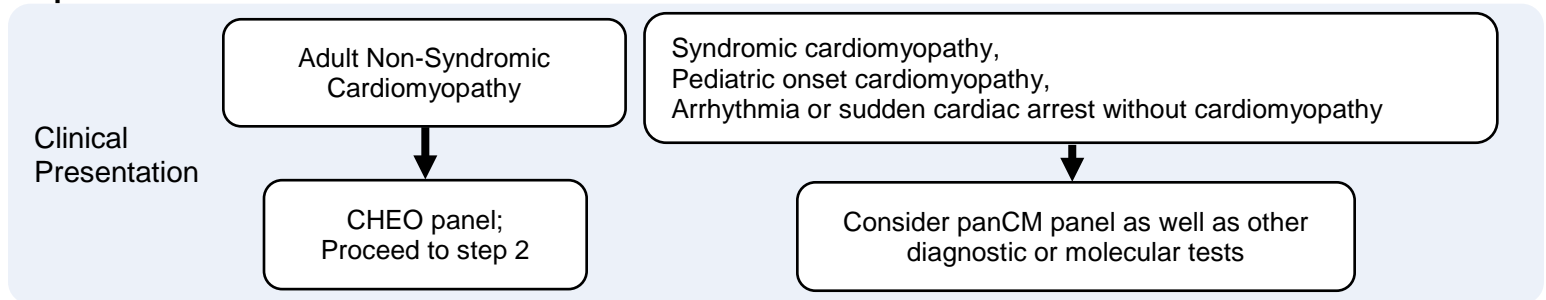
Genes included in panel: *DSC2*, *DSG2*, *DSP*, *FLNC*, *JUP*, *PKP2*, and *TMEM43* (c.1073C>T mutation only)  
 Analysis includes sequencing as described above, and MLPA of *PKP2*.

### Pan Cardiomyopathy Panel

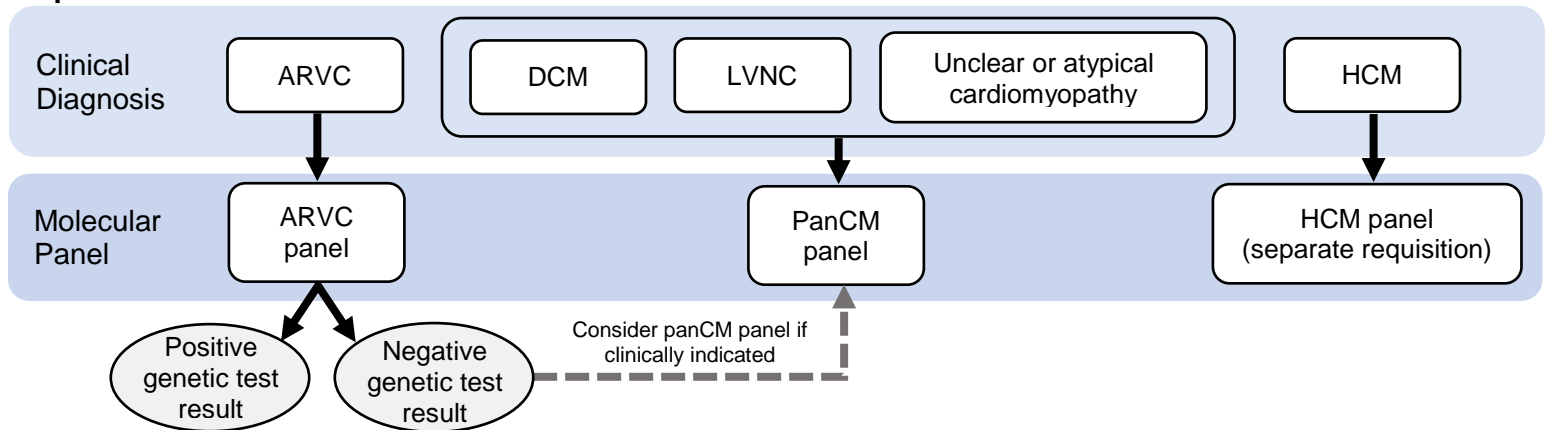
Genes included in panel: *ACTC1*, *ACTN2*, *BAG3*, *DES*, *DSC2*, *DSG2*, *DSP*, *FLNC*, *GLA*, *JUP*, *LAMP2*, *LMNA*, *MYBPC3*, *MYH7*, *MYL2*, *MYL3*, *NEXN*, *PKP2*, *PLN*, *PRKAG2*, *RBM20*, *SCN5A*, *TMEM43* (c.1073C>T mutation only), *TNNC1*, *TNNI3*, *TNNT2*, *TPM1*, *TTN*, *TTR*, and *VCL*  
 Analysis includes sequencing as described above, and MLPA of *BAG3*, *MYH7*, *MYBPC3*, *PKP2*, and *TNNT2*.

### Selecting a CHEO panel

#### Step 1:



#### Step 2:



### Other considerations:

- **LVNC**: If additional cardiac anomalies such as congenital heart disease, consider additional tests.
- **DCM**: If extra-cardiac signs, such as muscle weakness, hearing/vision loss or if arrhythmia > cardiomyopathy, consider other etiologies and tests
- **HCM**: In patients with a family history of non-HCM cardiomyopathy or sudden cardiac death, consider the panCM panel if indicated. Consider syndromic causes of HCM, particularly in young patients with severe disease. If there is uncertainty as to which panel to order or there are additional cardiac anomalies or family history, please speak with the lab genetic counsellor.