



## Clinical Policy: Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia

Reference Number: HNCA.CP.MP.493

Effective Date: 10/09

Last Review Date: 06/19

[Coding Implications](#)

[Revision Log](#)

See [Important Reminder](#) at the end of this policy for important regulatory and legal information.

### Description

Catecholaminergic polymorphic ventricular tachycardia (CPVT), also known as familial CPVT, is a rare disorder characterized by episodic syncope occurring during exercise or acute emotion in individuals without structural cardiac abnormalities and an unremarkable resting electrocardiogram (ECG).

### Policy/Criteria

- I. It is the policy of Health Net of California that genetic testing for CPVT is **medically necessary** for any of the following indications:
  - A. To confirm the diagnosis of CPVT in individuals who demonstrate exercise induced ventricular arrhythmias in the presence of an unremarkable ECG and absence of cardiac abnormalities
  - B. Individuals with a positive family history of sudden death in children or young adults and based on clinical evaluation (eg. history and physical, resting ECG, ECG and exercise stress testing), CPVT is suspected
  - C. Predictive testing for at-risk symptomatic family members when there is a positive genetic test in a first degree relative (i.e., individual's parents, full siblings, and children)

### Background

CPVT typically begins in childhood or adolescence, although cases have been reported with initial presentation in the fourth decade of life. Affected individuals may have a family history of juvenile sudden death or stress-induced syncope. CPVT may also present sporadically as a de novo mutation in individuals with no family history. Affected patients typically present with life threatening ventricular tachycardia or ventricular fibrillation occurring during emotional or physical stress, with syncope often being the first manifestation of the disease.

The two genes with mutations that have been identified with CPVT are the cardiac ryanodine receptor gene (which may have the p.G357S RyR2 mutation), an autosomal dominant form and the calsequestrin 2 gene, autosomal recessive. Both mutations appear to act by inducing diastolic calcium release from the sarcoplasmic reticulum. The resulting intracellular calcium overload leads to delayed afterdepolarizations and triggered activity, which can induce ventricular tachycardia and fibrillation. It should be noted that mutations in these two genes have been recognized in only 60% of patients with CPVT, implying that other genes may play a role. <sup>2</sup>

*American College of Cardiology/American Heart Association Task Force and the European Society of Cardiology*



## CLINICAL POLICY

### Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia

Beta blockers are recommended as first-line treatment in all affected individuals, (class I indication) and the use of the implantable cardioverter-defibrillator is recommended for higher-risk subgroups. The guidelines note that genetic analysis may help identify silent carriers of CPVT related mutations. Once identified silent carriers may be treated with beta blockers to reduce the risk of cardiac events and may receive appropriate genetic counseling to assess the risk of transmitting the disease to offspring. <sup>10</sup>

Most of the data available for inherited arrhythmogenic diseases such as Long QT syndrome, Brugada syndrome and CPVT, are derived from large registries that have followed patients over time, recording outcome information. They note that no randomized studies are available, and most likely they will never be conducted in these uncommon conditions. <sup>10</sup>

A study was completed to try to identify the genetic cause of CPVT in specific families, to preventively treat and clinically characterize the mutation-positive individuals, and to functionally characterize the pathogenic mechanisms of the mutation. Genetic testing was performed for 1404 relatives. Functional studies showed that the G357S mutation increased caffeine sensitivity and store overload-induced calcium release activity under conditions that mimic catecholaminergic stress. The study supports the use of genetic testing to identify individuals at risk of SCD to undertake prophylactic interventions. <sup>9</sup>

A number of other studies also note the importance of genetic testing for the early diagnosis of asymptomatic carriers, the prevention of sudden death, as well as to initiate appropriate prophylactic interventions. <sup>5689</sup>

### Coding Implications

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| CPT® Codes | Description   |
|------------|---|
| 81405      | Molecular Pathology Procedure, Level 6 (eg., analysis of 6-10 exons by DNA sequence analysis, mutation scanning or duplication /deletion variants of 11-25 exons) [i.e., CASQ2 (Calsequestrin 2, cardiac muscle), e.g., catecholaminergic polymorphic ventricular tachycardia] full gene sequence |



**CLINICAL POLICY**

**Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia**

| CPT® Codes | Description   |
|------------|---|
| 81408      | Molecular Pathology Procedure, Level 9, e.g., analysis of >50 exons in a single gene by DNA sequence analysis [i.e., RyR2 (ryanodine receptor 2 (cardiac)) e.g. catecholaminergic polymorphic Ventricular tachycardia, arrhythmogenic right ventricular dysplasia, full gene sequence of targeted sequence analysis >50 exons |
| 81413      | Cardiac ion channelopathies (eg, Brugada syndrome, long QT syndrome, short QT syndrome, catecholaminergic polymorphic ventricular tachycardia); genomic sequence analysis panel, must include sequencing of at least <u>10</u> genes, including ANK2, CASQ2, CAV3, KCNE1, KCNE2, KCNH2, KCNJ2, KCNQ1, RYR2, and SCN5A         |
| 81414      | Cardiac ion channelopathies (eg, Brugada syndrome, long QT syndrome, short QT syndrome, catecholaminergic polymorphic ventricular tachycardia); duplication/deletion gene analysis panel, must include analysis of at least 2 genes, including KCNH2 and KCNQ1  |

| HCPCS Codes | Description |
|-------------|-------------|
| N/A         |             |

**ICD-10-CM Diagnosis Codes that Support Coverage Criteria**

| ICD-10-CM Code | Description   |
|----------------|---|
| I47.2          | Ventricular tachycardia   |
| I49.041        | Ventricular fibrillation  |
| R55            | Syncope and collapse  |
| Z13.6          | Encounter for screening for cardiovascular disorders                |
| Z13.79         | Encounter for other screening for genetic and chromosomal anomalies |
| Z82.41         | Family history of sudden cardiac death                              |

| Reviews, Revisions, and Approvals  | Date  | Approval Date |
|--|-------|---------------|
| Policy Adopted from Health Net NMP # 493 Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia | 06/17 |               |
| Update and no changes  | 6/18  |               |

**References**

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## CLINICAL POLICY

### Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia

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### **Important Reminder**

This clinical policy has been developed by appropriately experienced and licensed health care professionals based on a review and consideration of currently available generally accepted standards of medical practice; peer-reviewed medical literature; government agency/program



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## CLINICAL POLICY

### Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia

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## CLINICAL POLICY

### Genetic Testing for Catecholaminergic Polymorphic Ventricular Tachycardia

**Note: For Medicaid members,** when state Medicaid coverage provisions conflict with the coverage provisions in this clinical policy, state Medicaid coverage provisions take precedence. Please refer to the state Medicaid manual for any coverage provisions pertaining to this clinical policy.

**Note: For Medicare members,** to ensure consistency with the Medicare National Coverage Determinations (NCD) and Local Coverage Determinations (LCD), all applicable NCDs, LCDs, and Medicare Coverage Articles should be reviewed prior to applying the criteria set forth in this clinical policy. Refer to the CMS website at <http://www.cms.gov> for additional information.

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