

# **Test Definition: DRPL**

## Dentatorubral-Pallidoluysian Atrophy (DRPLA) Gene Analysis, Varies

#### **Reporting Title:** DRPLA Gene Analysis **Performing Location:** Rochester

### Shipping Instructions:

Specimen preferred to arrive within 96 hours of collection.

#### **Specimen Requirements:**

Specimen Type: Whole blood

**Patient Preparation:** A previous bone marrow transplant from an allogenic donor will interfere with testing. Call 800-533-1710 for instructions for testing patients who have received a bone marrow transplant.

#### **Container/Tube:**

**Preferred:** Lavender top (EDTA) or yellow top (ACD)

Acceptable: Any anticoagulant

Specimen Volume: 3 mL

#### **Collection Instructions:**

1. Invert several times to mix blood.

2. Send whole blood specimen in original tube. Do not aliquot.

#### Forms:

1. New York Clients-Informed consent is required. Document on the request form or electronic order that a copy is on file. The following documents are available:

-Informed Consent for Genetic Testing (T576)

-Informed Consent for Genetic Testing (Spanish) (T826)

2. Molecular Genetics: Neurology Patient Information

3. If not ordering electronically, complete, print, and send a <u>Neurology Specialty Testing Client Test Request</u> (T732) with the specimen.

Specimen Type	Temperature	Time	Special Container	
Varies	Ambient (preferred)			
	Frozen			
	Refrigerated			

#### **Result Codes:**

Result ID	Reporting Name	Туре	Unit	LOINC®
53234	Result Summary	Alphanumeric		50397-9
53235	Result	Alphanumeric		49631-5
53236	Interpretation	Alphanumeric		69047-9
53237	Specimen	Alphanumeric		31208-2
53238	Source	Alphanumeric		31208-2
53239	Released By	Alphanumeric		18771-6

 $\ensuremath{\mathsf{LOINC}}\xspace^{\ensuremath{\mathbb{R}}}$  and CPT codes are provided by the performing laboratory.



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#### Supplemental Report:

No

### **CPT Code Information:**

81177-ATN1 (ataxin 2) (eg, denatatorubral-pallidolyuysian atrophy) gene analysis, evaluation to detect abnormal (eg, expanded) alleles

#### **Reference Values:**

Normal alleles: 7-35 CAG repeats Abnormal alleles: 49-93 CAG repeats

An interpretive report will be provided.