# **PRONTO<sup>®</sup> Tay-Sachs Ash kit**

## For the detection of the following mutations in Ashkenazi Jews in the HEXA gene:

1278insTATC IVS12+1G>C G269S (805G>A)

Instructions for Use



Tests





### INTENDED USE

The Pronto<sup>®</sup> Tay-Sachs Ash kit is a single nucleotide primer extension ELISA procedure, intended for the qualitative *in vitro* detection of the following three mutations in the HEXA gene c.1278insTATC, c.IVS12+1G>C and p.G269S in amplified human DNA.

For in vitro diagnostic use.

### BACKGROUND

Tay-Sachs disease is a fatal, autosomal recessive, neurodegenerative disorder caused by mutations in the HEXA gene, located on the human 15q23-q24 chromosome. It is characterized by normal motor development in the first few months of life followed by progressive weakness and loss of motor function, starting at 2-6 months of age, decreased social interaction, seizures and blindness, leading to total debilitation. Death typically occurs between 2 and 5 years of age. Tay-Sachs disease is caused by excess storage of GM2 ganglioside in lysosomes due to a deficiency of the isoenzyme hexosaminidase (hexA).

The carrier frequency for mutations that cause Tay-Sachs varies with ethnicity. The carrier frequency in Ashkenazi Jews is approximately 1 in 30.

Three mutations in the HEXA gene account for approximately 98% of all Tay-Sachs mutations found in the Ashkenazi Jewish population: c.1278insTATC 81%, c.IVS12+1G>C 15%, and p.G269S 2%. The overall carrier frequency for Tay-Sachs disease in non-Jewish individuals is 1 in 300, with a higher carrier rate in French-Canadian or Cajun descent.

In addition to general population screening, Tay-Sachs DNA testing is strongly recommended for individuals with ambiguous or positive HEXA enzyme test results.

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### REFERENCES

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- Kaback M, Lim-Steele J, Dabholkar D, Brown D, Levy N, Zeiger K. Tay-Sachs disease-carrier screening, prenatal diagnosis, and the molecular era. An international perspective, 1970 to 1993. The International TSD Data Collection Network. *J.A.M.A.*, 270:2307-15 (1993)
- 3. Kaback MM. Population-based genetics screening for reproductive counseling: the Tay-Sachs experience. *Eur. J. Peds.* 159:S192-5 (2000)
- 4. Navon R, Argov Z, Frisch A. Hexosaminidase A deficiency in adults. *Am. J. Med. Genet.*, 24:179-96 (1986)
- 5. American College of Obstetricians and Gynecologists (ACOG) Committee on Genetics. Pub. no. 162 (1995)
- 6. Okada S and O'Brien JS. Tay-Sachs disease: generalized absence of a beta-D-N- acetylhexosaminidase component. *Science*, 165:698-700 (1969)

### WARNINGS & PRECAUTIONS

- Reagents supplied in this kit may contain up to 0.1% sodium azide that is toxic if swallowed. Sodium azide has been reported to form explosive lead or copper azides in laboratory plumbing. To prevent the accumulation of these compounds, flush the sink and plumbing with large quantities of water.
- TMB Substrate solution is an irritant of the skin and mucous membranes. Avoid direct contact.
- In addition to reagents in this kit, the user may come in contact with other harmful chemicals that are not provided, such as ethidium bromide and EDTA. The appropriate manufacturers' Material Safety Data Sheets (MSDS) should be consulted prior to the use of these compounds.

### SASAY OVERVIEW

The PRONTO<sup>®</sup> procedure detects predefined polymorphisms in DNA sequences, using a single nucleotide primer extension ELISA procedure.

**1 TARGET DNA AMPLIFICATION:** The DNA fragments that encompass the tested mutations are amplified. This amplified DNA is the substrate for the primer extension reaction.

- 2 **POST-AMPLIFICATION TREATMENT:** The amplified DNA is treated to inactivate free unincorporated nucleotides, so that they will not interfere with the primer extension reaction.
- **3 PRIMER EXTENSION REACTION:** A single-nucleotide primer extension reaction is carried out in a 96-well thermoplate. Each well contains a 5'-labeled primer that hybridizes to the tested DNA next to the suspected mutation site, and a single biotinylated nucleotide species, which complements the nucleotide base at the tested site. Each post-amplification treated sample is tested in two wells per mutation: the first well of each pair tests for the presence of the mutant allele (*mut*), while the second well tests for the presence of the normal allele (*wt*). The biotinylated nucleotide will be incorporated to the primer in the course of the reaction or not added, depending on the tested individual's genotype.
- 4 **DETECTION BY ELISA:** The detection of the biotinylated primers is carried out by an ELISA procedure: The biotin-labeled primers bind to a streptavidin-coated ELISA plate and are detected by a peroxidase-labeled antibody (HRP) directed to the 5' antigenic moiety of the primer. A peroxidase reaction occurs in the presence of TMB-Substrate.
- 5 **INTERPRETATION OF THE RESULTS:** The results are determined visually (substrate remains clear or turns blue) or colorimetrically using an ELISA Reader.

### Isclaimer

- Results obtained using this kit should be confirmed by an alternative method.
- Confirmed results should be used and interpreted only in the context of the overall clinical picture. The manufacturer is not responsible for any clinical decisions that are taken.

The user of this kit should emphasize these points when reporting results to the diagnosing clinician or the genetic counselor.

### CONTENTS OF THE KIT

Tay-Sachs Ash Amplification Mix 1 vial	(0.95mL)
ProntoTaq™1 vial	(20 μL)
PRONTO <sup>®</sup> Buffer 2 1 bottle	e (3 mL)
Solution C 1 vial (	yellow cap) (130 μL)
Solution D 1 vial (	red cap) (100 μL)
ColoRed™ Oil 1 drop	per bottle (13 mL)
Assay Solution 1 bottle	e (green solution) (100 mL)
Wash Solution (conc. 20x) 1 bottle	e (100 mL)
HRP Conjugate 1 vial	(450 μL)
TMB Substrate 1 bottle	e (40 mL)
PRONTO <sup>®</sup> Tay-Sachs Ash Plates 3 indivi	dually pouched plates
Detection Plates 3 Strep	tavidin-coated ELISA plates

### STORAGE AND STABILITY

- Store the ProntoTaq<sup>™</sup> at -20°C.
- Do not freeze the kit.
- Do not use the kit beyond its expiration date (marked on box label). Stability is maintained even when components are re-opened several times.
- Minimize the time reagents spend at room temperature.
- This kit has been calibrated and tested as a unit; do not mix reagents from kits with different lot numbers.

### ADDITIONAL MATERIALS REQUIRED

- Deionized water (about two liters per kit)
- Thermowell plate or tubes (thin wall) for the post-amplification treatment
- Sterile pipette tips
- Troughs/reagent reservoirs for use with the detection reagents
- Thermocycler for a 96-well microplate
- Multichannel pipettes (5-50 µL and 50-200 µL)

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- Positive displacement pipettes (1-5  $\mu L,$  5-50  $\mu L,$  50-200  $\mu L$  & 200-1,000  $\mu L)$
- Filtered tips
- ELISA reader with 620 nm filter
- Polaroid camera and color film to record results (optional)
- Automated microtiter plate washer or squirt bottle
- Vortex mixer
- Timer

### S ASSAY PROCEDURE

### **1.** DNA AMPLIFICATION

- 1. **Dispense** 2  $\mu$ L template DNA (from an initial concentration of about 150 ng/ $\mu$ L) to a Thermowell plate or tube.
- 2. **Prepare** a Master Mix in a sterile vial, according to the volumes indicated in the table below, plus one spare reaction volume. Add the ProntoTaq<sup>™</sup> to the Master Mix shortly before dispensing the mix. Gently mix by pipetting in and out several times.

### PCR Master mix

Solution	Volume for one sample
Amplification Mix Tay-Sachs Ash	13.0 μL
ProntoTaq™	0.2 μL

- 3. Dispense 13 µL Master Mix to each well or tube.
- 4. Add one drop of ColoRed<sup>™</sup>-Oil to each well. Do not touch the wells with the tip of the oil bottle. Even when using a thermocycler with a hot lid, it is recommended to use oil.

5. **Place** the Thermowell plate or tube in a thermocycler previously programmed with the following protocol:

### Cycling protocol

1.	94°C	5 minutes
2. 3. 4.	94°C 63°C 72°C	30 seconds 30 seconds 30 seconds 35 cycles
5.	72°C	5 minutes

6. To verify amplification, **subject** 5 μL of the amplified product to electrophoresis in a 2% agarose gel.

#### Sizes of amplified fragments:

Gene	Mutation	Fragment size
HEXA	c.1278insTATC c.IVS12+1G>C	600 bp
	p.G269S	293 bp

#### Limitation of the test:

Different thermocyclers may influence the amplification yield dramatically. It is recommended to use a calibrated thermocycler.

### **2** POST- AMPLIFICATION TREATMENT

Only 5  $\mu$ L of each amplified DNA sample will be used to carry out this assay.

1 **Prepare** a post-amplification treatment mix shortly before use. Combine in a single test tube the volumes appearing in the following table, multiplied by the number of tested samples, plus one spare volume.

#### **Post-Amplification Mix**

Solution	Volume for one sample
PRONTO <sup>®</sup> Buffer 2	47.0 µL
Solution C	2.0 µL
Solution D	1.5 µL

- 2 Mix gently by pipetting this solution in and out five times. Do not vortex.
- 3 Add 50  $\mu$ L of the post-amplification mix into each Thermowell or tube containing 5  $\mu$ L of each amplified DNA sample.

Ensure that the solution you add becomes well mixed with the DNA sample by pipetting.

- 4 Add one drop of ColoRed<sup>™</sup> oil to each tube. Do not touch the tube with the tip of the oil bottle. Even when using a thermocycler with a hot lid, it is essential to use oil.
- **5 Incubate** for 30 minutes at 37°C, then for 10 minutes at 95°C in a thermocycler.

If not used immediately, the treated sample can be kept at 2-8°C for a maximum of four hours.

### **3** PRIMER EXTENSION REACTION

**1 Program** the thermocycler as follows:

Cycle	Temperature	Time
Start:	94°C	15 sec.
20 cycles:	{ 94°C 62°C	30 sec.
20 090103.	ι <sub>62°C</sub>	10 sec.
End:	18-25°C - Coo	I down to room temperature

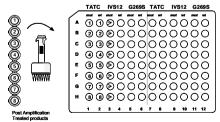
**2 Take** a PRONTO<sup>®</sup> Plate out of its pouch. Notice the color at the bottom of the wells. For each mutation tested, use a pink well (*mut*) and a blue well (*wt*). Mark the plate with the ID numbers of your test.

If you intend to use less than a full plate, you can cut the plate and return the unused portion to the pouch. If you do this, seal the pouch immediately with its desiccant card inside.

**3** Dispense 8 μL of post-amplification treated DNA into the first six wells in row A (see Fig. 1). Continue with the remaining samples. It is possible to transfer up to eight samples simultaneously using a multichannel pipette.

Ensure that the solution is at the bottom of each well by inspecting the plate from below. Make sure that the well does not contain air bubbles.

Figure 1: Scheme for dispensing Post-Amplification Treated DNA samples into the PRONTO<sup>®</sup> Tay-Sachs Ash Plate



### **Recommendation:**

Use a new set of tips for each column. Alternatively use the same set of tips, but do not touch the bottom of the wells.

- 4 Tilt the plate and add one drop of ColoRed<sup>™</sup> Oil to each well. Do not touch the well with the tip of the oil bottle. Even when using a thermocycler with a hot lid, it is not essential to use oil.
- **5 Turn on** the thermocycler and start the cycling protocol.
- **6** When the thermal cycling is complete, you can proceed to the ELISA, or store the reaction products in the refrigerator and carry out the visualization steps within 24 hours.

### 4 ELISA ASSAY - COLOR DEVELOPMENT

The ELISA consists of the following steps:

- Binding the biotinylated primer to the Streptavidin-coated plate.
- Washing away the unbound primer.
- Incubating with the HRP conjugate.
- Washing away the unbound conjugate.
- Incubating with the TMB Substrate (color development).

The results of this assay can be determined in one of two ways:

**a Visually**: by monitoring the development of the blue color.

or

**b** Colorimetrically: by measuring the absorbance using an ELISA reader at a wavelength of 620 nm.

### PREPARATION

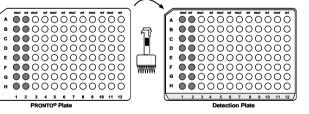
- All components used in the detection step should reach room temperature before starting the assay.
- Dilute the 20x Wash Solution to 1x with deionized water.
   Dilute solution may be kept at 18-25° C for up to one month.
- **Peel** off the plastic cover of the Detection Plate. Mark the side of the plate with the kit name and test number.
- Place the PRONTO<sup>®</sup> Plate and the Detection Plate side by side, oriented in the same direction (see Fig. 2).

### DETECTION BY ELISA

- **1 Fill** a reagent reservoir /trough with the green colored Assay Solution. About 11 mL will be required for a 96-well plate.
- 2 Using a multichannel pipette, add 100 μL of Assay Solution to the bottom of each well in column 1 of the PRONTO<sup>®</sup> Plate. Gently mix by pipetting in and out 3-4 times.
- **3** Without changing tips, transfer 100 µL from each well in this column to the first column in the Detection Plate (see Fig. 2).

Ensure that the solution at the bottom of all wells of the PRONTO<sup>®</sup> plate has turned green by inspecting them from below.

## Figure 2: Transferring the primer extension products from the PRONTO<sup>®</sup> Plate to Detection Plate.



**4 Repeat** this procedure, using a new set of tips for each column. It is essential to maintain the order of the samples.

10  $\mu$ L of oil carried over or 10  $\mu$ L of the sample left behind will not significantly affect the detection process.

5 Incubate for 10 minutes at room temperature (18-25°C).

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- **6** While the incubation takes place, dilute the Conjugated HRP 1:100 in assay solution (green solution). About 11 mL are needed for a 96-well plate. This solution must be freshly prepared each time the test is run.
- **7 Empty** the plates, wash four times with 350 μL 1x Wash Solution. Ensure that the plates are dry after the last wash step.
- **8** With a multichannel pipette add 100 μL freshly-diluted Conjugated HRP to all the wells.
- 9 Incubate for 10 minutes at RT.
- **10 Wash** as in step 7.
- **11** Add 100 μL TMB substrate to each well with a multichannel pipette and incubate for 15 minutes at RT (18-25°C) until the blue color appears sufficiently strong.
- **12 For Visual Detection:** Results may be documented by a standard Polaroid camera with color film (for example, Fuji FP-100C).
- **13 For Colorimetric Detection: Agitate** the plate gently to homogenize the color in the wells. **Read** the results in an ELISA reader using a 620 nm filter (singe wavelength setting).

### VALIDATION OF THE RESULTS

#### For Visual Detection:

For every mutation site tested, at least one of the wells should develop a deep blue color. Otherwise, the results are invalid for the relevant mutation (see Fig. 3).

### For Colorimetric Detection:

For every mutation site tested, at least one of the two wells should yield an O.D.  $\ge$  0.50 reading.

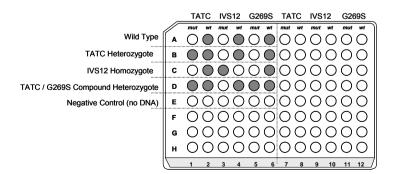
### INTERPRETATION OF RESULTS

**Important:** Heterozygous or homozygous mutant results should be confirmed by retesting. It is recommended to repeat the test with newly extracted DNA.

### **Criteria for Visual Interpretation**

A deep blue color indicates a positive signal, while negative signals appear as a clear or pale blue-colored well (see Fig. 3).

## Figure 3: Genotype assignment according to visual inspection of test results



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### Criteria for Colorimetric Interpretation

The genotype of each sample is determined according to two criteria:

- 1. The O.D. values of the *mut* and *wt* wells.
- 2. The ratio of *mut / wt* O.D. values.

Calculate the *mut / wt* ratios by dividing the signal of the *mut* well by the signal of the *wt* well.

Identify the correct genotype using the table below:

Genotype	<i>mut</i> well	wt well	mut/wt ratio
Normal	O.D. <u>&lt;</u> 0.35	O.D. <u>≥</u> 0.5	ratio ≤0.5
Heterozygote	O.D. <u>≥</u> 0.5	O.D. <u>≥</u> 0.5	0.5 < ratio < 2.0
Homozygote	O.D. <u>≥</u> 0.5	O.D. <u>&lt;</u> 0.35	ratio ≥ 2.0

Samples with values not included in the above table are considered indeterminate and should be retested.

### PRONTO® Tay-Sachs Ash - PROCEDURE SUMMARY

DNA EXTRACTION: from human wh	olo blood using a validated	mothod
	ole blood, using a validated i	
DNA AMPLIFICATION:		
Volumes per reaction: 2 µL template	e DNA + 13 µL Amplification I	Mix + 0.2 µL
ProntoTaq™		
Cycling protocol:	/ <u>22 <sup>0</sup>2 22 / <del>7</del>2 <sup>0</sup>2 22</u>	) <del>7</del> 0%0 5 ·
94°C 5 min $\rightarrow$ 35 cycles of {94 °C 30 set	ec. / 63 °C 30 sec. / 72 °C 30	sec.} $\rightarrow$ 72 °C 5 min.
POST-AMPLIFICATION PROCEDUR		
<ul> <li>Volumes per reaction:</li> </ul>	PRONTO <sup>®</sup> Buffer 2	47.0 μL
	Solution C	2.0 µL
	Solution D	1.5 µL
Pipette in and out to mix.		
<ul> <li>Add 50 µL into each well containin</li> </ul>	a 5 ul amplified product mix	x well
<ul> <li>Add so µ2 into each wer containing</li> <li>Add one drop of ColoRed™ oil.</li> </ul>		wen.
<ul> <li>Incubate 30 minutes at 37° C, ther</li> </ul>	10 minutes at 95°C	
PRIMER EXTENSION REACTION:		
<ul> <li>Dispense 8 µL of each Post-Amp</li> </ul>	lification treated DNA into si	x wells of the PRONTO <sup>®</sup>
Plate.		
<ul> <li>Add one drop of ColoRed<sup>™</sup> oil.</li> </ul>		
Start the cycling protocol:		
94°C 15 sec→20 cycles of {94°C	30 sec. / 62°C 10 sec.} →Co	ool.
Insert the PRONTO <sup>®</sup> Plate in the therm	nocycler when the temperatu	re is 90°C.
DETECTION:		
Add 100 µL Assay Solution to each	well in the PRONTO® plate a	and mix.
Transfer 100 µL from each well of t	he PRONTO <sup>®</sup> Plate to the re	espective position in the
Detection Plate. Incubate 10 minute	es at RT.	
Empty the wells and wash four time	•	
Add 100 µL 1:100 Conjugate HRP 1	to every well; incubate for 10	minutes at RT.
<ul> <li>Wash the wells again.</li> </ul>		
<ul> <li>Add 100 µL Substrate to each well;</li> </ul>	incubate at RT for 15 minute	es.

For a troubleshooting guide, please refer to our website: <u>www.prontodiagnostics.com/ts</u>

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The Pronto<sup>®</sup> Technology is covered by US patent 5,710,028, by European patent 0648222 and by corresponding national patents.

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