

NIEMANN-PICK TYPE B ΔR608 TEST KIT

This kit provides the components needed to detect the Niemann-Pick type B polymorphism 1821-1823 3 bases deletion in the Sphingomyelin Phosphodiesterase 1 (SMPD1) gene (Chromosome 11p15.4-p15.1) resulting in an Arg608 amino acid deletion.

Store at -20°C

For in vitro use only

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1. PRINCIPLE

Heteroduplex Analysis Test Kits contain DUPLEXER™ reagents. These are synthetic oligonucleotides that mimic naturally occurring gene segments of human DNA, but their structures may be manipulated and defined.

When amplified gene segments derived from human genomic DNA and the DUPLEXER™ reagent are mixed the resultant heteroduplexes possess unique 3-D structures specific to a particular genetic allele.

The genotype of clinical samples may be determined by comparison of the electrophoretic mobility patterns of heteroduplex bands from test DNA with control DNAs provided with the kit.

There are three main stages involved in distinguishing between wild homozygous, mutant homozygous and heterozygous alleles;

1. Amplification of Control DNAs, DUPLEXER™ and test genomic DNA in a PCR reaction.
2. Crossmatching of the DUPLEXER™ PCR product with the PCR products from the control and test DNAs.
3. Separation of the resulting heteroduplexes by polyacrylamide gel electrophoresis or other appropriate acrylamide based electrophoretic separation technique.

The principle of genotyping using DUPLEXER™ reagents is fully outlined on the Heteroduplex Analysis web site. It is recommended that the user studies the principle and examples provided on the web pages if unfamiliar with DUPLEXER™ technology.

CAUTION

DUPLEXER™ reagent is more easily amplified than genomic DNA. To minimise cross contamination ALWAYS open the DUPLEXER™ tube after completing the set-up of the rest of your PCR reaction tubes.

2. KIT CONTENT

The Niemann-Pick Δ R608 Test Kit contains sufficient reagents for 10 tests.

| | |
|------------|-----------------|
| Clear cap | 'Buffer' |
| White cap | 'R608 Test' |
| Green cap | 'R608 DUPLEXER' |
| Red cap | 'R608 Wild' |
| Yellow cap | 'R608 Mutant' |

Kits in their lyophilized form may be stored for up to 6 months at -20°C.

Reconstitution of Reagents

Prior to use the lyophilized reagents must be reconstituted by addition of buffer solution as follows;

| | |
|---------------|------------------------|
| R608 Test | Add 264 μ l Buffer |
| R608 DUPLEXER | Add 336 μ l Buffer |
| R608 Wild | Add 336 μ l Buffer |
| R608 Mutant | Add 336 μ l Buffer |

The contents of the tubes should be thoroughly mixed by vortexing (10 seconds) and quick spinning at least three times.

Reconstituted reagents may be stored for up to 3 months at -20°C.

NOTE: It is recommended that stored reagents are mixed as above after being thawed completely.

Reagent to be supplied by user

Taq DNA polymerase (0.5 units/ μ l)

3. DNA SAMPLE PREPARATION

This kit is intended for use with genomic DNA. Samples prepared using commercial DNA purification kits or by phenol/chloroform extraction may be used.

It is recommended that DNA is suspended in DNase-free water although the use of buffers supplied with kits will have no adverse effect.

Samples should be stored at 4°C or at -20°C for longer periods.

Genomic DNA should be diluted to a concentration of 10-100ng/μl before use.

4. PROCEDURE

STEP 1: PCR reaction

| Tube | Mix for 1 PCR reaction (μl) | | | |
|--------------------|-----------------------------|---------------------|----------|------------|
| | Wild Control CGC | Mutant Control ΔCGC | Test DNA | DUPLEXER * |
| Red cap | 28 | - | - | - |
| Yellow cap | - | 28 | - | - |
| White cap | - | - | 22 | - |
| Patient DNA | - | - | 6 | - |
| Green cap | - | - | - | 28 |
| Taq (0.5 units/μl) | 2 | 2 | 2 | 2 |

*To run a complete set of controls (including heterozygotes) plus:

1 to 2 DNA tests use 1 DUPLEXER reaction tube.

3 to 7 DNA tests use 2 DUPLEXER reaction tubes.

8 to 10 DNA tests use 3 DUPLEXER reaction tubes.

Thermal Cycling Parameters;

Initial Denaturation (94°C, 5 min) – 1 cycle

Denaturation (94°C, 30 sec)
Primer annealing (64°C, 30 sec)
Primer extension (72°C, 30 sec) } 32 cycles

Final primer extension (72°C, 5 min) – 1 cycle

If the PCR products are not to be processed immediately store at -20°C.



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STEP 2: 'Crossmatching' - Formation of Heteroduplexes

In this step, aliquots of the test DNA and control DNA PCR products are mixed with aliquots of the DUPLEXER™ PCR product. The mixtures are placed in a thermal cycler and subjected to heating and cooling to induce the formation of heteroduplexes, a process referred to as crossmatching. Note: the control heterozygote (genotype CGC/ΔCGC) is generated by mixing Wild control CGC (genotype CGC/CGC) and mutant control ΔCGC (genotype ΔCGC/ΔCGC) PCR products in equal proportions.

To ensure optimal heteroduplex formation, approximately equal molar amounts of PCR product from the DUPLEXER™ and test DNAs should be used in the crossmatch mixtures. Therefore before crossmatching we recommend testing of each test DNA, control DNA, and DUPLEXER™ PCR product to compare band intensities. This can be easily achieved by electrophoresis of 5µl of each DNA in 15% polyacrylamide gels or 2% agarose minigels.

Adjustment of the PCR product concentration can be made based on visual comparison of the band intensity.

Mix contents of the PCR tubes by briefly vortexing and centrifuging and prepare crossmatch reactions.

| PCR Product | Mix for 1 Crossmatch reaction (µl) | | | |
|-----------------------|------------------------------------|--------------|-------------------|----------|
| | Control CGC | Control ΔCGC | Control CGC/ ΔCGC | Test DNA |
| DUPLEXER | 5 | 5 | 5 | 5 |
| Control CGC (wild) | 5 | - | 5 | - |
| Control ΔCGC (mutant) | - | 5 | 5 | - |
| Test DNA | - | - | - | 5 |

Thermal Cycling Parameters;

Increase temperature to 95°C, hold for 2 minutes
Cool to 60°C at 1°C per second, hold for 1 minute
Cool to 45°C at 0.1°C per second, hold for 1 minute
Cool to 6°C

If the crossmatched products are not to be processed immediately store at -20°C.

STEP 3: Electrophoresis of Heteroduplexes

Electrophoresis may be performed in a range of electrophoresis equipment:

Large format gel (Nominal length 16cm)

- 15% (37.5:1) non-denaturing polyacrylamide gel
- 1x TBE running buffer
- Constant voltage of 500V
- Gel cooling to maintain a temperature of 10-15°C. Cold tap water cooling is sufficient

Midi format gel (Nominal length 9cm)

- 15% (37.5:1) non-denaturing polyacrylamide gel
- 1.5x TBE running buffer
- Constant voltage of 300V
- Gel cooling to maintain a temperature of 10-15°C. Cold tap water cooling is sufficient

An appropriate molecular weight marker, for example a 100 base pair ladder, should be included.

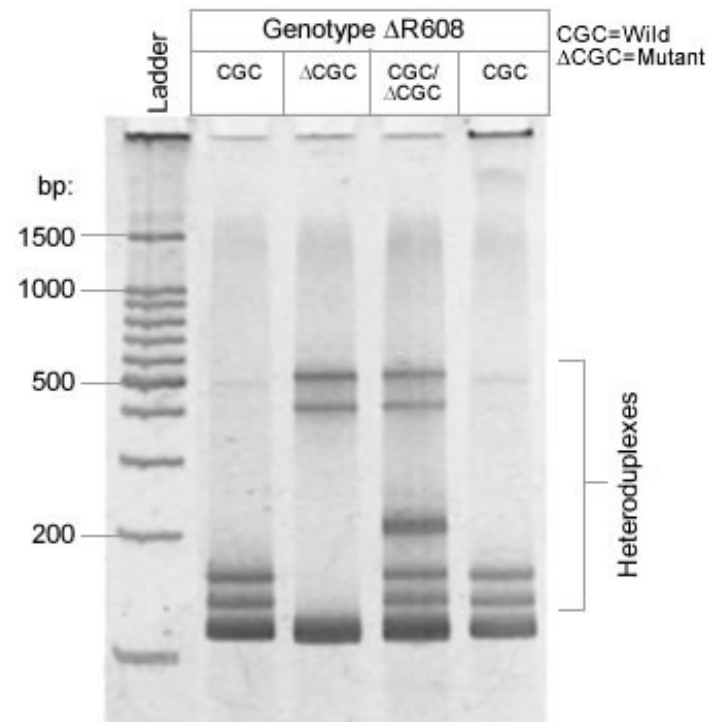
An appropriate amount of gel loading buffer should be added to each crossmatch reaction before loading the complete reaction mixture onto the gel. We recommend a gel loading buffer containing xylene cyanol.

For good resolution of heteroduplexes gels should be run until the xylene cyanol marker reaches the bottom of the gel.

Gels may be stained using ethidium bromide or SYBR Green.

NOTE: Heteroduplex Analysis kits may also be used in conjunction with most other acrylamide-electrophoretic based separation techniques.

STEP 4: Analysis of the Stained Gel



Genotypes of clinical samples may be determined by comparison of electrophoretic mobility patterns of heteroduplex bands with the control genomic DNAs provided with the kit.

The patterns are;

- Homozygous wild type DNA (CGC)
- Homozygous mutated or polymorphic DNA (Δ CGC)
- Heterozygous wild type plus mutated or polymorphic DNA (CGC/ Δ CGC)
- Homozygous wild type DNA (CGC)

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