DNA Test Results (form D)

Please complete the survey below.

Thank you!

D1.0 - If you have the DNA Test Report, please scan and upload the complete report.

D1.1 - Was a genetic exam performed?

○ Yes
○ No
(if No, please skip DNA Test Results survey.)

D1.2 - Where was the genetic test done?

_________________________

D1.3 - Is a sample of the DNA stored in the lab?

○ Yes
○ No

D1.4 - YEAR of the genetic test?

_________________________
(type in year only (e.g. 2017))
Gene Searching and Mutations/Variations

Report 1

D2.1 - Is this gene mutation associated with MH or Central Core Disease?
○ Yes
○ No

D2.2 - Gene Name
○ RYR1
○ CACNA1S
○ STAC3
○ other (specify)

D2.2.1 - If other, please specify:
__________________________________

D2.3 - Gene symbol
(type in gene symbol, ref. OMIM (e.g. RYR1, ARG2458His))

D2.4 - the Exon #:
__________________________________

D2.5 - Aminoacid change
(e.g. p.2458Arg>His)

D2.6 - Nucleotide change
(e.g. c.488G>T)
Gene Searching and Mutations/Variations

Report 2

D3.1 - Is this mutation associated with MH or Central Core Disease?
○ Yes
○ No

D3.2 - Gene Name
○ RYR1
○ CACNA1S
○ STAC3
○ other (specify)

D3.2.1 - If other, please specify:
__________________________________

D3.3 - Gene symbol
(type in gene symbol, ref. OMIM (e.g. RYR1, ARG2458HIS))

D3.4 - the Exon #:
__________________________________

D3.5 - Aminoacid change
(e.g. p.2458Arg>His)

D3.6 - Nucleotide change
(e.g. c.488G>T)
## Gene Searching and Mutations/Variations

### Report 3

<table>
<thead>
<tr>
<th>D4.1 - Is this mutation associated with MH or Central Core Disease?</th>
<th>Yes</th>
<th>No</th>
</tr>
</thead>
<tbody>
<tr>
<td>D4.2 - Gene Name</td>
<td><img src="#" alt="Options" /></td>
<td></td>
</tr>
<tr>
<td>D4.2.1 - If other, please specify:</td>
<td><img src="#" alt="Other Gene" /></td>
<td></td>
</tr>
<tr>
<td>D4.3 - Gene symbol</td>
<td><img src="#" alt="Gene Symbol" /></td>
<td></td>
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<tr>
<td>D4.4 - the Exon #:</td>
<td><img src="#" alt="Exon Number" /></td>
<td></td>
</tr>
<tr>
<td>D4.5 - Aminoacid change</td>
<td><img src="#" alt="Aminoacid Change" /></td>
<td></td>
</tr>
<tr>
<td>D4.6 - Nucleotide change</td>
<td><img src="#" alt="Nucleotide Change" /></td>
<td></td>
</tr>
</tbody>
</table>

- **Yes**: RYR1, CACNA1S, STAC3, other (specify)
- **No**: 

### Gene Options

- **RYR1**
- **CACNA1S**
- **STAC3**
- **other (specify)**

### Gene Symbol Details

- **(type in gene symbol, ref. OMIM (e.g. RYR1, ARG2458His))**

### Exon Number

- **(e.g. p.2458Arg>His)***

### Aminoacid Change

- **(e.g. c.488G>T)***
Other Gene Mutations/Variations:

(If you need more space to report additional gene mutations/variations, please contact NAMHR administrator. We will send you the instruction and format how you can submit your complete genetic report.)

D5.1 - Were any other GENE sequence variants?

☐ Yes
☐ No

D5.1.1 - If yes, please report with the following order:

(GENE_NAME, GENE_SYMBOL, EXON#, AMINOACID_CHANGE, NUCLEOTIDE_CHANGE)

D5.2 - please list all other BENIGN variants not yet included above, with the following order:

(GENE_NAME, GENE_SYMBOL, EXON#, AMINOACID_CHANGE, NUCLEOTIDE_CHANGE)

!This field is for official use only!
NEXT SURVEY

Do you still want to fill out the Clinical Information about this participant?

- Yes
- No