Usher Syndrome Type 3A

Usher 3A is a rare genetic disorder. It is characterized by hearing and vision loss that begins in late childhood and worsens over time. A person must have two variants in the CLRN1 gene in order to have this condition.

Erin, you **do not have the variant** we tested.

You could still have a variant not covered by this test.

How To Use This Test

This test does not diagnose any health conditions.

Please talk to a healthcare professional if this condition runs in your family, you think you might have this condition, or you have any concerns about your results.

- Review the Carrier Status tutorial
- See Scientific Details

Intended Uses

- To test for the N48K variant in the CLRN1 gene.
- To identify carrier status for Usher 3A.

Limitations

- Does not test for all possible variants for the condition.
- Does not report if someone has two copies of a tested variant.
- Does not cover other subtypes of Usher syndrome.

Important Ethnicities

- This test is most relevant for people of Ashkenazi Jewish descent.
- This test does not include variants commonly found in people of Finnish descent with Usher 3A.

You are likely not a carrier.

This result is relevant for you because you have Ashkenazi Jewish ancestry.

We ruled out the most common variant for Usher 3A in people of Ashkenazi Jewish descent.

You still have a chance of being a carrier for Usher 3A.

You may still have up to a **1 in 1,700 chance** of carrying a variant not covered by this test.

See Scientific Details
About Usher Syndrome Type 3A

When symptoms develop
Symptoms typically develop during late childhood or adolescence.

How it’s treated
There is currently no known cure. Hearing loss may be treated with hearing aids. Vision loss may be monitored with routine eye exams. Early intervention is recommended to teach alternative communication skills.

Typical signs and symptoms
- Hearing loss in childhood or early teens
- Gradual vision loss
- Night blindness by mid-teens
- Blindness by mid-adulthood

Ethnicities most affected
This syndrome is most common in people of Ashkenazi Jewish and Finnish descent.

Read more at
Genetics Home Reference
National Institute on Deafness and Other Communication Disorders

Consider talking to a healthcare professional if you are concerned about your results.

If you’re starting a family, a genetic counselor can help you and your partner understand if additional testing might be appropriate.

Connect with a GIC

Share your results with a healthcare professional.

Print report

Learn more about this condition and connect with support groups.

Learn more

Usher 3A is caused by variants in the CLRN1 gene.

The CLRN1 gene contains instructions for making a protein called clarin 1. This protein is found in the ears and eyes, although its exact function is unknown. Certain variants in CLRN1 result in a protein with the wrong shape that does not function properly.

Read more at Genetics Home Reference
You have no variants detected by this test.

<table>
<thead>
<tr>
<th>Marker Tested</th>
<th>Your Genotype*</th>
<th>Additional Information</th>
</tr>
</thead>
<tbody>
<tr>
<td>N48K</td>
<td>A</td>
<td>Biological explanation</td>
</tr>
<tr>
<td>Gene: CLRN1</td>
<td>Typical copy from your other parent</td>
<td>Typical vs. variant DNA sequence(s)</td>
</tr>
<tr>
<td>Marker: N490151</td>
<td></td>
<td>Percent of 23andMe customers with variant</td>
</tr>
</tbody>
</table>

*This test cannot distinguish which copy you received from which parent. This test also cannot determine whether multiple variants, if detected, were inherited from only one parent or from both parents. This may impact how these variants are passed down.

23andMe always reports genotypes based on the “positive” strand of the human genome reference sequence (build 37). Other sources sometimes report genotypes using the opposite strand.

Test Interpretation

Post-Test Carrier Risk

This report provides an estimate of the post-test carrier risk for people of Ashkenazi Jewish descent only.

- For people with partial ethnicity from one or more groups mentioned above, post-test carrier risk depends on the exact mixture in the person’s background.
- Post-test risk for other ethnicities cannot be provided because sufficient data is not available.

<table>
<thead>
<tr>
<th>Post-test carrier risk for relevant ethnicities</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Ashkenazi Jewish</td>
<td>1 in 1,700</td>
</tr>
</tbody>
</table>
Indications for Use

The 23andMe PGS Carrier Status Test for Usher Syndrome Type 3A is indicated for the detection of the NM8K variant in the CIRN1 gene. This test is intended to be used to determine carrier status for Usher 3A in adults, but cannot determine if a person has two copies of a tested variant. The test is most relevant for people of Ashkenazi Jewish descent.

Special Considerations

• There are currently no professional guidelines in the U.S. for carrier testing for this condition.

Test Performance Summary

Carrier Detection Rate & Relevant Ethnicities

The "carrier detection rate" is an estimate of the percentage of carriers for this condition that would be identified by this test. Carrier detection rate differs by ethnicity and is provided only where sufficient data is available.

<table>
<thead>
<tr>
<th>Ethnicity</th>
<th>Rate (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ashkenazi Jewish</td>
<td>93%</td>
</tr>
<tr>
<td>Finnish</td>
<td></td>
</tr>
</tbody>
</table>

Analytical Performance

Accuracy was determined by comparing results from this test with results from sequencing for 49 samples with known variant status. 49 out of 49 genotype results were correct. Fewer than 1 in 100,000 samples may receive a Not Determined result. This can be caused by random test error or unexpected DNA sequences that interfere with the test. It can also be caused by having two copies of a variant tested.
Warnings and Limitations

- This test does not cover all variants that could cause this condition.*
- This test does not diagnose any health conditions.
- Positive results in individuals whose ethnicities are not commonly associated with this condition may be incorrect. Individuals in this situation should consider genetic counseling and follow-up testing.
- Share results with your healthcare professional for any medical purposes.
- If you are concerned about your results, consult with a healthcare professional.

See the Package Insert for more details on use and performance of this test.

* Variants not included in this test may be very rare, may not be available on our genotyping platform, or may not pass our testing standards.

References


