Hereditary Fructose Intolerance

Hereditary fructose intolerance is a rare genetic disorder. It is characterized by low blood sugar levels, stomach pain, and vomiting after eating fructose. A person must have two variants in the ALDOB gene in order to have this condition.

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

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

0 variants detected in the ALDOB gene

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

- Tests for multiple variants in the ALDOB gene.
- To identify carrier status for hereditary fructose intolerance.

Limitations

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

Important Ethnicities

- This test is most relevant for people of European descent.

You are likely not a carrier.

This result is relevant for you because you have European ancestry.

We ruled out the most common variants for hereditary fructose intolerance in people of European descent.

You still have a chance of being a carrier for hereditary fructose intolerance.

You may still have up to a 1 in 240 chance of carrying a variant not covered by this test.

See Scientific Details
### About Hereditary Fructose Intolerance

#### When symptoms develop
Symptoms typically develop during infancy.

#### How it’s treated
There is currently no known cure. Maintaining a fructose-free diet may reduce or prevent symptoms.

#### Typical signs and symptoms
- Nausea and vomiting
- Low blood sugar
- Stomach pain
- Failure to gain weight
- Liver disease
- Kidney disease

#### Ethnicities most affected
This condition occurs in people of all ethnicities, but is best studied in people of European descent.

**Read more at**
- Genetics Home Reference
- MedlinePlus
- National Organization for Rare Disorders

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**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 GC](#)

Share your results with a healthcare professional.

[Print report](#)

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**Hereditary fructose intolerance is caused by variants in the ALDOB gene.**

The ALDOB gene contains instructions for making an enzyme called aldolase B. This enzyme is found mainly in the liver, where it helps to break down the sugar fructose. Certain variants in ALDOB disrupt this function, resulting in a harmful buildup of fructose byproducts in liver cells.

**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>A149P</td>
<td>C</td>
<td>&gt; Biological explanation</td>
</tr>
<tr>
<td>Gene: ALDOB</td>
<td>Typical copy from one of your parents</td>
<td>&gt; Typical vs. variant DNA sequence(s)</td>
</tr>
<tr>
<td>Marker: rs1800566</td>
<td>C</td>
<td>&gt; Percent of 23andMe customers with variant</td>
</tr>
<tr>
<td></td>
<td>Typical copy from your other parent</td>
<td>&gt; References [5, 6, 7, 8, 10, 11]</td>
</tr>
<tr>
<td>N334K</td>
<td>G</td>
<td>&gt; Biological explanation</td>
</tr>
<tr>
<td>Gene: ALDOB</td>
<td>Typical copy from one of your parents</td>
<td>&gt; Typical vs. variant DNA sequence(s)</td>
</tr>
<tr>
<td>Marker: rs12665</td>
<td>G</td>
<td>&gt; Percent of 23andMe customers with variant</td>
</tr>
<tr>
<td></td>
<td>Typical copy from your other parent</td>
<td>&gt; References [3, 5, 6, 7, 8, 9, 11]</td>
</tr>
<tr>
<td>Delta4E4</td>
<td>TTTG</td>
<td>&gt; Biological explanation</td>
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<td>&gt; References [1, 4, 7, 8, 11]</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 European descent only.

- For people of partial European descent, post-test carrier risk is less than that for those who are fully European. The exact post-test risk depends on how much European ancestry a person has.
- Post-test risk for other ethnicities cannot be provided because sufficient data is not available.

Post-test carrier risk for relevant ethnicities

<table>
<thead>
<tr>
<th>Ethnicity</th>
<th>Risk</th>
</tr>
</thead>
<tbody>
<tr>
<td>European</td>
<td>1 in 240</td>
</tr>
</tbody>
</table>
Indications for Use

The 23andMe PGS Carrier Status Test for Hereditary Fructose Intolerance is indicated for the detection of three variants in the ALDOB gene. This test is intended to be used to determine carrier status for hereditary fructose intolerance in adults, but cannot determine if a person has two copies of a tested variant. The test is most relevant for people of European 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>European</td>
<td>70%</td>
</tr>
</tbody>
</table>

Analytical Performance

Accuracy was determined by comparing results from this test with results from sequencing for 149 samples with known variant status. 149 out of 149 genotype results were correct. Fewer than 1 in 100,000 samples may receive a Not Determined result for one or more variants included in this test. 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


