

PEX1 gene

peroxisomal biogenesis factor 1

Normal Function

The *PEX1* gene provides instructions for making a protein called peroxisomal biogenesis factor 1 (Pex1p), which is part of a group of proteins called peroxins. Peroxins are essential for the formation and normal functioning of cell structures called peroxisomes. Peroxisomes are sac-like compartments that contain enzymes needed to break down many different substances, including fatty acids and certain toxic compounds. They are also important for the production of fats (lipids) used in digestion and in the nervous system. Peroxins assist in the formation (biogenesis) of peroxisomes by producing the membrane that separates the peroxisome from the rest of the cell and by importing enzymes into the peroxisome. Pex1p enables other peroxins to bring enzymes into the peroxisome.

Health Conditions Related to Genetic Changes

Zellweger spectrum disorder

At least 114 mutations in the *PEX1* gene have been identified in people with Zellweger spectrum disorder, which is a group of conditions that have overlapping signs and symptoms and affect many parts of the body. The conditions' features, which vary in severity, can include weak muscle tone (hypotonia), developmental delay, and vision and hearing problems. Mutations in the *PEX1* gene are the most common cause of Zellweger spectrum disorder and are found in nearly 70 percent of affected individuals.

There are two common *PEX1* gene mutations found in people with Zellweger spectrum disorder. One mutation replaces the protein building block (amino acid) glycine with the amino acid aspartic acid at position 843 in Pex1p (written as Gly843Asp or G843D). This mutation leads to reduced levels of the protein. Individuals who have the G843D mutation tend to have signs and symptoms that are at the less-severe end of the condition spectrum. The other common mutation, which is known as the 1700fs mutation, leads to the production of an abnormally short, nonfunctional Pex1p. People who have the 1700fs mutation often have signs and symptoms that are at the severe end of the condition spectrum.

Mutations in the *PEX1* gene that cause Zellweger spectrum disorder reduce or eliminate the activity of the Pex1p protein. Without enough functional Pex1p, enzymes are not

properly imported into peroxisomes. As a result, cells contain empty peroxisomes that cannot carry out their usual functions. The severe end of the condition spectrum is caused by the absence of functional peroxisomes within cells. The less severe end of the condition spectrum results from mutations that allow some peroxisomes to form.

Other Names for This Gene

- peroxin1
- peroxisome biogenesis disorder protein 1
- PEX1_HUMAN
- Pex1p
- ZWS1

Additional Information & Resources

Tests Listed in the Genetic Testing Registry

- Tests of PEX1 ([https://www.ncbi.nlm.nih.gov/gtr/all/tests/?term=5189\[geneid\]](https://www.ncbi.nlm.nih.gov/gtr/all/tests/?term=5189[geneid]))

Scientific Articles on PubMed

- PubMed (<https://pubmed.ncbi.nlm.nih.gov/?term=%28%28PEX1%5BTIAB%5D%29+OR+%28peroxin1%5BALL%5D%29+OR+%28Pex1p%5BALL%5D%29%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+3600+days%22%5Bdp%5D%29>)

Catalog of Genes and Diseases from OMIM

- PEROXISOME BIOGENESIS FACTOR 1; PEX1 (<https://omim.org/entry/602136>)

Gene and Variant Databases

- NCBI Gene (<https://www.ncbi.nlm.nih.gov/gene/5189>)
- ClinVar ([https://www.ncbi.nlm.nih.gov/clinvar?term=PEX1\[gene\]](https://www.ncbi.nlm.nih.gov/clinvar?term=PEX1[gene]))

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Genomic Location

The *PEX1* gene is found on chromosome 7 (<https://medlineplus.gov/genetics/chromosome/7/>).

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