



NOD2 gene

nucleotide binding oligomerization domain containing 2

Normal Function

The *NOD2* gene (previously known as *CARD15*) provides instructions for making a protein that plays an important role in immune system function. The NOD2 protein is active in some types of immune system cells (including monocytes, macrophages, and dendritic cells), which help protect the body against foreign invaders such as bacteria and viruses. The protein is also active in several types of epithelial cells, including Paneth cells, which are found in the lining of the intestine. These cells help defend the intestinal wall against bacterial infection.

The NOD2 protein has several critical functions in defending the body against foreign invaders. The protein is involved in recognizing certain bacteria and stimulating the immune system to respond appropriately. When triggered by specific substances produced by bacteria, the NOD2 protein turns on (activates) a protein complex called nuclear factor-kappa-B. This protein complex regulates the activity of multiple genes, including genes that control immune responses and inflammatory reactions. An inflammatory reaction occurs when the immune system sends signaling molecules and white blood cells to a site of injury or disease to fight microbial invaders and facilitate tissue repair.

The NOD2 protein also appears to play a role in a process called autophagy, which cells use to surround and destroy bacteria, viruses, and other harmful substances. In addition to protecting cells from infection, autophagy is used to recycle worn-out cell parts and break down certain proteins when they are no longer needed. This process is also involved in the self-destruction of cells (apoptosis).

Health Conditions Related to Genetic Changes

Blau syndrome

At least 22 mutations in the *NOD2* gene have been found to cause Blau syndrome, an inflammatory disorder that begins in childhood and primarily affects the skin, joints, and eyes. These mutations change single protein building blocks (amino acids) in the NOD2 protein. All of these mutations result in a version of the NOD2 protein that is overactive, which can trigger an abnormal inflammatory reaction and cause swelling and irritation. However, it is unclear how the abnormally active protein causes the specific pattern of inflammation affecting the skin, joints, and eyes that is characteristic of Blau syndrome.

NOD2 gene mutations can also cause early-onset sarcoidosis, a similar condition that some researchers consider to be a noninherited version of Blau syndrome.

Crohn disease

Variations in the *NOD2* gene have been associated with an increased risk of Crohn disease, a complex disorder that causes inflammation of the digestive system. In particular, *NOD2* gene changes are associated with a form of Crohn disease that affects the lower part of the small intestine (the ileum) and the colon in populations of northern European descent. The three most common *NOD2* variations are found in about 40 percent of all people with Crohn disease. The most common of these, written as 3020insC or 1007fs, leads to the production of a *NOD2* protein that is slightly shorter than normal. Other common variations change single amino acids in the *NOD2* protein. It is unclear how these genetic changes increase the risk of developing Crohn disease. Studies suggest that changes in the *NOD2* gene prevent the protein from recognizing bacteria appropriately, allowing these microbes to grow unchecked and invade cells that line the intestine. An abnormal immune response to these bacteria may contribute to inflammation and the digestive problems characteristic of Crohn disease.

Yao syndrome

Several variations in the *NOD2* gene increase the risk of developing Yao syndrome, a disorder that causes episodes of fever and abnormal inflammation affecting many parts of the body. Most people with this condition have a variation in the *NOD2* gene written as IVS8+158, which is located in a region of the gene called intron 8. Another relatively common *NOD2* gene variation in people with Yao syndrome replaces the amino acid arginine with the amino acid tryptophan at position 702 in the *NOD2* protein (written as Arg702Trp or R702W). These variations have also been identified in some people with Crohn disease (described above). About 20 percent of people with Yao syndrome have both the IVS8+158 and R702W variations, and some affected individuals have additional variations in the *NOD2* gene.

The effects of having one, two, or more variations in the *NOD2* gene is unclear, although studies suggest that these genetic changes may alter the activity (expression) of the *NOD2* gene. It is unknown how these variations might contribute to abnormal inflammation in people with Yao syndrome. Researchers suspect that environmental factors such as infections may also play a role in triggering the disease in people with genetic variants that increase their risk.

Cancers

A few studies have suggested a possible association between changes in the *NOD2* gene, particularly the common variation 3020insC (described above), and the development of several types of cancer. Although some of these studies found an increased risk of cancer in people with a *NOD2* gene variation, other research found no such association. It is unclear how changes in this gene might contribute to cancer risk.

Other disorders

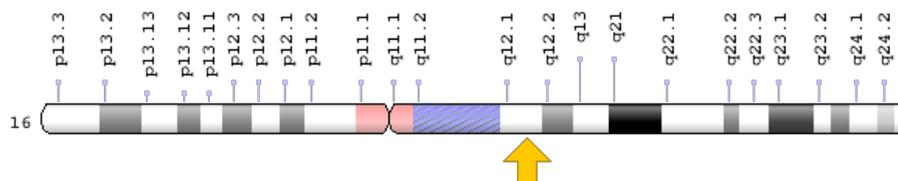
Several studies have considered variations in the *NOD2* gene as a possible risk factor for a condition called graft-versus-host disease (GVHD). GVHD can occur following certain cancer treatments, such as allogeneic stem cell transplantation. This procedure, which is typically used to treat cancers of the blood and immune system, replaces a patient's diseased blood-forming cells (a type of stem cell) with stem cells from a healthy donor. If the donor's stem cells (the graft) recognize the patient's body (the host) as foreign, they may attack organs and tissues such as the liver, digestive system, and skin. GVHD is the term used to describe this potentially severe reaction.

A few studies have suggested that variations in the *NOD2* gene influence the risk of developing severe complications of GVHD following an allogeneic stem cell transplant. The presence of *NOD2* gene variations in both the stem cell donor and the recipient is associated with the greatest risk of a severe reaction. However, other research has found no relationship between *NOD2* gene changes and the risk of developing GVHD.

Chromosomal Location

Cytogenetic Location: 16q12.1, which is the long (q) arm of chromosome 16 at position 12.1

Molecular Location: base pairs 50,693,587 to 50,733,077 on chromosome 16 (Homo sapiens Annotation Release 109, GRCh38.p12) (NCBI)



Credit: Genome Decoration Page/NCBI

Other Names for This Gene

- ACUG
- BLAU
- CARD15
- caspase recruitment domain family, member 15
- caspase recruitment domain protein 15
- CD
- IBD1

- inflammatory bowel disease protein 1
- LRR-containing protein
- NOD2_HUMAN
- NOD2B
- nucleotide-binding oligomerization domain containing 2
- PSORAS1

Additional Information & Resources

Educational Resources

- Immunobiology: The Immune System in Health and Disease (fifth edition, 2001):
The Front Line of Host Defense
<https://www.ncbi.nlm.nih.gov/books/NBK27105/>

Scientific Articles on PubMed

- PubMed
<https://www.ncbi.nlm.nih.gov/pubmed?term=%28%28CARD15%5BTIAB%5D%29+OR+%28NOD2%5BTIAB%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+360+days%22%5Bdp%5D>

Catalog of Genes and Diseases from OMIM

- NUCLEOTIDE-BINDING OLIGOMERIZATION DOMAIN PROTEIN 2
<http://omim.org/entry/605956>

Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology
http://atlasgeneticsoncology.org/Genes/GC_NOD2.html
- ClinVar
<https://www.ncbi.nlm.nih.gov/clinvar?term=NOD2%5Bgene%5D>
- HGNC Gene Symbol Report
https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:5331
- Monarch Initiative
<https://monarchinitiative.org/gene/NCBIGene:64127>
- NCBI Gene
<https://www.ncbi.nlm.nih.gov/gene/64127>
- UniProt
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Reviewed: December 2017
Published: June 11, 2019

Lister Hill National Center for Biomedical Communications
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